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GenCore version 5.1.6

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OM protein - protein search, using sw model

Run on: December 28, 2005, 13:43:04; Search time 147.538 Seconds (without alignments)
375.236 Million cell updates/sec
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126 S .....NYYYYGMDVWGQGTTVTVS residues QVQLVESGGGVVQPGRSLRL... 439378781 Gapext 0.5 US-09-920-137F-7 668 2443163 seqs, BLOSUM62 Gapop 10.0 , score: table: Sequence: Searched: Title: Perfect Scoring

Total number of hits satisfying chosen parameters: 2443163
Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database : A\_Geneseq\_21:\*

1: geneseqp1980s:\*
2: geneseqp2000s:\*
4: geneseqp2001s:\*
5: geneseqp2001s:\*
7: geneseqp2003as:\*
7: geneseqp2003bs:\*
8: geneseqp2004s:\*
9: geneseqp2005s:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

1		<b>*</b>			SUMMARIES	
No.	OL	Match	Length	DB	ID	Description
! ! ! +-! ! !	657.5		146	5	AAM51171	Aam51171 H
~	52.	7.76	146	Ŋ	AAM51170	Aam51170 Human rec
m	27.	93.9	4	Ŋ	AAM51169	
4	20.	92.9	146	ហ	AAM51172	I
ഹ	15.	N	146	വ	AAM51168	168
9	58		135	7	ADD28320	Add28320 Human het
7	æ	7	135	7	ADD28236	36
80	æ	7	135	σ	ADV86823	$\sim$
σ	æ	87.1	124	9	ADA89238	Ada89238 Human ant
	81.	7.	127	~	AAY17954	4
	•	7.	ሞ	7	AAW80815	Aaw80815 Amino aci
	81.	7	457	σ	4	-4
	81.	7	523	ო	AAY44994	4
	81.	7.	524	ო	AAY44995	995
	æ	7	126	7	ADK18593	
	œ	7.	126	7	ADK18785	Adk18785 Anti-huma
	œ	7.	126	7	ADK18858	Adk18858 Anti-huma
	₿	7.	126	œ	ADL25404	
	7	9	126	7	ADK18786	98
	7	9	126	7	ADK18822	Adk18822 Anti-huma
	7	9	126	7	ADK18882	2 Anti-
	-	9	126	7	ADK18601	Adk18601 Anti-huma
	7	9	126	œ	ADL25420	Ad125420 Human mAb
24	~	9	126	æ	ADP22222	Adp22222 Human ant

25 8 ADP22316 41 9 ADV67314 18 5 AAM51167 25 8 ADP22300 25 8 ADP22312 34 8 ADP22312 52 7 ADG96547 51 3 AAY93734 51 3 AAY93734 51 6 AAE35889 26 7 ADC28172 25 7 ADC2817 25 7 ADE28451 27 ADE28475	2316 Human a 1164 Anti-tu 7314 Amino a 1167 Human D 2300 Human a 2312 Human a 2348 Human a	45720 Human B 96547 Single 93734 The hea 35889 Human 1	dk18821 dk884372 dr68514 dd28317	Ade28451 Human ant Aeb27731 Prostate Aeb27751 Anti-pros Ade28475 Human ant Aeb45960 Human mon
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77.5 86. 577 86. 577 86. 575 86. 75.5 86. 72.5 86. 72.5 85. 72.5 85. 72.5 85. 71.5 85. 71.5 85. 71.5 85. 71.5 85.	3335 330 330 330 330 330	W W W W V	4 3 3 4 4 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	4 4 4 4 4 4 4 4 4 4 1 4 6 4 6

## ALIGNMENTS

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Tumour necrosis factor alpha; TNF; antibody; heavy chain; CDR; complementarity determining region; antirheumatic; antiarthritic; antiulcer; antiasthmatic; antiallergic; antiinflammatory; antisickling; antidiabetic; antiarteriosclerotic; antiatherosclerotic; vasotropic; antianginal; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy; TNV148; monoclonal antibody; mAb.
                                                                                                                                                                                                                                 may
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                                                                                                                                                                                                                         , are PCR primer-encoded
sequence"
                                                                          chain variable region.
                                                                                                                                                                                                                                                                                                                                                                               Ser"
                                                                                                                                                                                                                                                                                                                                                                               substituted by
                                                                                                                                                                                                             1. .19
/label= Signal_peptide
/note= "amino acids 1-7
differ from the native a
                                                                                                                                                                                                                                                              'label= Mature_protein
                                                                                                                                                                                                                                                                                                                                                                               note= "wild-type Pro
                                                                          mAb TNV148(B) heavy
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                                                                                                                                                                                                    Location/Qualifiers
                  AAM51171 standard; protein; 146 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                     note= "encoded
                                                                                                                                                                                                                                                                                                                                       /label= CDR2
86. .117
/label= FR3
                                                                                                                                                                                                                                                                                                                                                                                                  CDR3
                                                                                                                                                                                                                                                                                          50. .54
/label= CDR1
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/label= FR2
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                                                                                                                                                                                                                                                                                                                                                                                                                              146
= J6
                                                        (first entry)
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/label=
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/label=
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/label=
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                                                                           Human recombinant
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Synthetic.
                                                        10-JUN-2002
                                     AAM51171;
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Peptide
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       AAM51171
RESULT
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07-AUG-2001;
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                                                                                      AAM51170
120
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AAM5117
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                                                                                                                                                                                                                                                                                               The present sequence is that of the heavy chain variable region of antitumour necrosis factor (TNF) human recombinant monoclonal antibody (mAb) TNV148 [8] was produced by site-directed mitagenesis of TNV148 (see AAM51170), replacing residue pro-94 with a germine Ser. TNV148 was (see AAM51170), replacing residue pro-94 with a germine Ser. TNV148 was to fise AAM51170), replacing residue pro-94 with a germine Ser. TNV148 was to fise AAM51170), replacing residue pro-94 with human constant region antibody transgenes that was immunised with human TNF alpha. The human mAbs bound immobilised human TNF alpha with high avidity and had a totally human 1971, kappa isotype. They showed relatences to each other and to the human germine DF-46 heavy chain sequence (see AAM51167). The invention and/or complementarity determining region (CDR)-grafted anti-TNF antibodies, immunoglobulins, clasvage products and other specified and/or complementary nucleic acids, vectors, host cells, compositions, formulations, devices, transgenic animals, transgenic compositions, formulations, devices, transgenic animals, the heavy chain and/or light chain variable regions given in the present sequence and in AAM51165, or aither all of the CDRs of the light chain (see AAM51161.

Sea AAM51156-60) or all of the CDRs of the light chain (see AAM51161.

Sea AAM51156-60) or all of the CDRs of the light chain (see AAM51161.

Sea AAM51156-60) or all of the CDRs of the light chain (see AAM51161.

Sea AAM51156-60) or all of the CDRs of the light chain (see AAM51161.

Sea AAM51156-60) or all of the CDRs of the light chain and condition in a cell, tissue, org
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYYGMDVWGQG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 146;
                                                                                                                                                          Shealy D;
                                                                                                                                                                                                                          antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                        mammalian anti-tumor necrosis factor anti
s cell anemia, diabetes, atherosclerosis,
s, myocardial infarction, leprosy.
                                                                                                                                                          Scallon B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ore 657.5; DB 5; ed. No. 7.3e-49; Mismatches 0;
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                                                                                                                                                           Heavner
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Pred.
                                                                                                                                                                                                                                                                             131pp; English
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                                                                            2000US-0223360P
2000US-0236826P
2001US-00920137
                                                   2001WO-US024785
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99.2%;
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treating sickle cell
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                                                                                                                                                                                                                                                                              4 ;
                                                                                                                                                                                                                                                    angina pectoris,
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                                                                                                                                                                                               N-PSDB; ABL53511
                                                                                                                                                                                                                                                                              3; Fig
WO200212502-A2
                                                                                                                                                          Giles-Komar J,
                                                                            07-AUG-2000;
29-SEP-2000;
01-AUG-2001;
                                                   07-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Sim:
Matches 126;
                          14-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               80
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  may
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                                                                                                                                                                                                                                                                                                                                                                                                            /label= Signal_peptide
/note= "amino acids 1-7 are PCR primer-encoded and
differ from the native sequence"
20. .146
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                                                                                                               standard; protein; 146
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2000US-0236826P.
2001US-00920137.
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/label= J6
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angina pectoris, myo
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126
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N-PSDB; ABL53510.
                                                                                                                                                                                                                 recombinant
TTVTVSS
                               TTVTVSS
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29-SEP-2000;
01-AUG-2001;
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tumour necrosis factor (TNF) human recombinant monoclonal antibody (mAb)
TNV148. TNV148 was 1 of 8 human mAbs produced from a GenTNV fusion using spleen cells from an electrical interval and constant region antibody transgenes that was immunised with human TNF alpha. The human mAbs bound immobilised human TNF alpha with high avidity and had a totally human igGl, kappa isotype. They showed relatedness to each other totally human germline DF-46 heavy chain sequence (see AAMSIL67). The invention provides isolated human, primate, rodent, mammalian, chimeric, humanised and/or complementarity determining region (CDR)-grafted anti-TNF antibodies, immunoglobulins, cleavage products and other specified portions and variants, as well as anti-TNF antibody compositions, formulations, devices, transgenic animals, transgenic plants, and methods of making and using them. The anti-TNF antibody compositions, formulations, devices, transgenic animals, transgenic plants, and methods of making and using them. The anti-TNF antibody compositions, formulations, dan immunoglobulin molecule, especially the heavy chain and/or light chain variable regions given in the present sequence and in AAMSIGS, or either all of the CDRs of the heavy chain (see AAMSIGS, or either all of the CDRs of the heavy chain sequence and in AAMSIGS, or either all of the CDRs of the heavy chain greatence and in AAMSIGS, or either all of the CDRs of the DRs of the heavy chain and/or light chain variable regions given in the present inhibit TNF induced cell adhesion molecules, inhibit TNF binding to receptor, or provide Arthritic Index improvement

C see AAMSILS8-60) or all of the CDRs of Arthritic Index improvement

C see AAMSILS9-60) or all of the CDRs of the light chain (see AAMSILS9-60) or all of the CDRs of the CDRs of an arteriosclerosis, atherosclerosis, atherosclerosis, atherosclerosis, atherosclerosis, atherosclerosis, atherosclerosis, atherosclerosis, atherosclerosis, atherosis, and fungal inferctions, pneumonia, leptosy and malaria, and subsemma, chron
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Sequence 146 AA;

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ij
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                                                                                                  139
                                       9
                                                          79
                                                                                         ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGN-YYYYGMDVWGQG
                                       DGSNKKY
                                                          20 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                    Gaps
                   1;
                                        QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSY
 Length 146;
                    Indels
 5
                    1;
Score 652.5; DB Pred. No. 2e-48;
           d. No. 2e-/
Mismatches
                    ;
0
97.7%;
llarity 98.4%;
Conservative
                                                                                                                                          146
Query Match
Best Local Similarity
                                                                                                                      TTVTVSS
                                                                                                                                         TTVTVSS
                  125;
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                    Matches
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Human recombinant mAb TNV15 heavy chain variable region.
          standard; protein; 146
                                 (first entry)
                                 10-JUN-2002
                      AAM51169;
          AAM51169
    AAM51169
RESULT
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Tumour necrosis factor alpha; TNF; antibody; heavy chain; CDR; complementarity determining region; antirheumatic; antiarthritic; antiulcer; antiasthmatic; antiallergic; antiinflammatory; antisickling; antidiabetic; antiarteriosclerotic; antiatherosclerotic; vasotropic; antianginal; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy; TNV15; monoclonal antibody; mAb.

sapiens Homo

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Location/Qualifiers
  Key
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Novel isolated mammalian anti-tumor necrosis factor antibody, useful for treating sickle cell anemia, diabetes, atherosclerosis, restenosis, angina pectoris, myocardial infarction, leprosy.
             may

    19
    /label= Signal_peptide
    /note= "amino acids 1-7 are PCR primer-encoded and differ from the native sequence"

                                                                                                                                                                                                                     Shealy D;
                                                                                                                                                                                                                    Scallon B,
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                               Mature protein
                                                                                                                    'note= "encoded by A"
                                                                                                                                                                                                                     Heavner
                                                                                                                                                                                2000US-0223360P.
2000US-0236826P.
2001US-00920137.
                                                                               CDR2
                                                                                                                                                                    2001WO-US024785
                                                      CDR1
                                                                                                        CDR3
                                                            55. .68
/label= FR2
                                           FR1
                                                                                           label= FR3
                                                                                                                         136. .146
/label= J6
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W
                                                                                                 118. .135
/label= CD
                        20. .146
/label= N
                                                                                     117
                                    31. .49
/label=
                                                                        /1abel=
                                                50. .54
/label=
                                                                                                                                                                                                                     Knight
                                                                                                  118.
                                                                                                                                                                                                         CENTOCOR INC
                                                                         69
                                                                                                                                                                                                                                 WPI; 2002-217194/27
                                                                                                                                                                                                                                       N-PSDB; ABL53509
                                                                                                             Misc-difference
                                                                                                                                                                                                                    ۵,
                                                                                                                                           WO200212502-A2
                                                                                                                                                                                07-AUG-2000;
29-SEP-2000;
                                                                                                                                                                                             01-AUG-2001;
                                                                                                                                                                    07-AUG-2001;
                                                                                                                                                                                                                     Giles-Komar
                                                                                                                                                        14-FEB-2002
                                                                                                                                                                                                         (CENZ)
Peptide
                         Protein
                                                                                     Region
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                                                Region
                                                             Region
                                                                         Region
                                                                                                   Region
                                                                                                                          Region
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4; 131pp; English. Example 3; Fig

The present sequence is that of the heavy chain variable region of antitumour necrosis factor (TNF) human recombinant monoclonal antibody (mAb) tumour necrosis factor (TNF) human recombinant monoclonal antibody (mAb) TNV15. TNV15 was 1 of 8 human mAbs produced from a GenTNV fusion using spleen cells from a hybrid mouse containing human variable and constant region antibody transgenes that was immunised with human TNF alpha. The human mAbs bound immobiles that was immunised with high avidity and had a totally human igG1, kappa isotype. Thy showed relatedness to each other and to the human germline DP-46 heavy chain sequence (see AAM51167). The invention provides isolated human, primate, rodent, mammalian, chimeric, humanised and/or complementarity determining region (CDR)-grafted anti-TNF antibodies, immunoglobulins, cleavage products and other specified portions and variants, as well as anti-TNF antibody compositions, formulations, devices, transgenic animals, transgenic plants, and methods of making and using them. The anti-TNF antibody compositions for making and using them. The anti-TNF antibody the heavy chain and/or light chain variable regions given in the present sequence and in AAM51165, or either all of the CDRs of the heavy chain (see AAM51161-63). The antibodies may inhibit TNF-induced cell adhesion molecules, inhibit TNF binding to receptor, or provide Arthritic Index improvement in a mouse model. They are useful for diagnosing or treating a TNF related condition in a cell, tissue, organ or animal (claimed) such as the context of the context of the claim or claims or the context of the claim of the claim of the claim or chain or claims. in a mouse model. They are useful for diagnosing or treating a TNF related condition in a cell, tissue, organ or animal (claimed) such as rheumatoid arthritis, gastric ulcer, asthma, allergic rhinitis, Crohn's pathology, sickle cell anaemia, diabetes, a cardiovascular disease such as arteriosclerosis, atherosclerosis, restenosis, angina pectoris or myocardial infarction, an infectious disease in a cell such as bacterial,

2001WO-US024785

07-AUG-2001;

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multiple
                                                                                                                                                                      119
                                                                                                                                                                                          139
 malignant
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                                                                                                                                                                                                                                                                                                                                                                      Tumour necrosis factor alpha; TNF; antibody; heavy chain; CDR; complementarity determining region; antirheumatic; antiarthritic; antiulcer; antiasthmatic; antiallergic; antiinflammatory; antisickling; antidiabetic; antiarteriosclerotic; antiatherosclerotic; vasotropic; antianginal; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy; TNV196; monoclonal antibody; mAb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and may
                                                                                                                                                                      ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARDRGIAAGGN-YYYYGMDVWGOG
                                                                                                                                              DGSNKKY
                                                                                                                             DGSNKKY
                                                                                                                                                                                         MDVWGQG
                                                                                                         Gaps
viral, and fungal infections, pneumonia, leprosy and malaria, a mal disease such as leukaemia, chronic myelocytic leukaemia, Burkitt's lymphoma and multiple myeloma, or a neurological disease such as musclerosis, Parkinson's disease, spinal ataxia, Alzheimer's disease Creutzfeldt-Jakob disease
                                                                                                          1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             are PCR primer-encoded
                                                                                                                                                                                 QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSY
                                                                                                                                         Length 146;
                                                                                                                                                                                                                                                                                                                                                    Human recombinant mAb TNV196 heavy chain variable region.
                                                                                                          Indels
                                                                                                         4.
                                                                                    5;
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/label= Signal peptide
/note= "amino acids 1-7 are PCR
differ from the native sequence
20. .146
                                                                                               .8e-46;
                                                                                    DB
                                                                                                        Mismatches
                                                                                    Score 627.5;
Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note= "encoded by
                                                                                                                                                                                                                                                                                       AAM51172 standard; protein; 146 AA
                                                                                                         3;
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                                                                                   93.9%;
illarity 93.7%;
Conservative
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/label= J6
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50
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/label=
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/label=
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/label=
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/label=
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                                                                                              Local Similarity
nes 119; Conser
                                                                                                                                                                                                                TTVTVSS
                                                                                                                                                                                                                                   TTVTVSS
                                                                 Sequence 146 AA;
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                                                                                    Query Match
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The present sequence is that of the heavy chain variable region of antitumour necrosis factor (TMF) human recombinant monoclonal antibody (mAb) tumour necrosis factor (TMF) human recombinant monoclonal antibody (mAb) produced from a definity tumon using spleen cells from a hybrid mouse containing human variable and constant region antibody transgenes that was immunised with human TMF alpha. The human mAbs bound immobilised human TMF alpha with high avidity and had a totally human IgG1, Kappa isotype. They showed relatedness to each other and to the human germline DP-46 heavy chain sequence (see AAM51167). The invention provides isolated human, primate, rodent, mammalian, chimeric, humanised and/or complementarity determining region (CDR)-grafted anti-TMF antibodies, immunoglobulins, cleavage products and other specified portions and variants, as well as anti-TMF antibody compositions, formulations, devices, transgenic animals, transgenic compositions, formulations, devices, transgenic animals, transgenic plants, and methods of making and using them. The anti-TMF antibody comprises at least a portion of an immunoglobulin molecule, especially the heavy chain and/or light chain variable regions given in the present sequence and in AAM51165 or either all of the CDRs of the heavy chain (see AAM51165) or all of the CDRs of the light chain (see AAM51161-63). The antibodies may inhibit TMF-induced cell adhesion molecules, inhibit TMF binding to receptor, or provide Arthritic Index improvement in a mouse model. They are useful for diagnosing or treating a TMF inhibit TMF planding or related condition in a cell, itssue, organ or animal (claimed) such as pathology, sickle cell anaemia, diabetes, a cardiovascular disease such as leutemenia, atherogenicular disease in a cell such as bacterial, wiral, and fungal infections, pneumonia, leprosy and malaria, a malignant consense, spinal ataxia, Alzebase such as leutemenia, a perion of provides and and multiple myeloma, or a neurological disease and such as leutemenia. Plymphoma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                QVQLVESGGGVVQPGRSLRLSCAASGFIPSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                               mammalian anti-tumor necrosis factor antibody, useful cell anemia, diabetes, atherosclerosis, restenosis, wyocardial infarction, leprosy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1;
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                                                                                                                                            Shealy
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                                                                                                                                            Scallon
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Pred. No. 1.1e-45
; Mismatches
                                                                                                                                            Ö
                                                                                                                                             Heavner
                                                                                                                                                                                                                                                                                                        131pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4;
                                 2000US-0223360P
2000US-0236826P
2001US-00920137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92.9%;
larity 92.9%;
Conservative
                                                                                                                                            Knight DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sclerosis, Parkinson's di
Creutzfeldt-Jakob disease
                                                                                                                                                                                                                                 Novel isolated mammal
treating sickle cell
angina pectoris, myoo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                126
                                                                                                         CENTOCOR INC
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hes 118; Conser
                                                                                                                                                                                                                                                                                                          4;
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                                                                                                                                                                                               ABL53512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 146 AA;
                                                                                                                                         ٦,
                                                                                                                                                                                                                                                                                                           3; Fig
                                 07-AUG-2000;
29-SEP-2000;
01-AUG-2001;
                                                                                                                                          Giles-Komar
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Best Local
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                                                                                                          (CENZ)
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RESULT 5

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nour necrosis factor alpha; TNF; antibody; heavy chain; CDR; plementarity determining region; antirheumatic; antiarthritic; iulcer; antiasthmatic; antiallergic; antiinflammatory; antisickling; idiabetic; antiarteriosclerotic; antiatherosclerotic; vasotropic; langinal; cardiant; antibacterial; virucide; fungicide; antileprotic; tozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; an; diagnosis; therapy; TNV14; monoclonal antibody; mAb.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is that of the heavy chain variable region of antitumour necrosis factor (TNF) human recombinant monoclonal antibody (mAb) TNV14. TNV14 was 1 of 8 human mAbs produced from a GenTNV fusion using spleen cells from a hybrid mouse containing human variable and constant region antibody transgenes that was immunised with human TNF alpha. The human mAbs bound immobilised human TNF alpha with high avidity and had totally human IgG1, kappa isotype. They showed relatedness to each other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        овів,
                                                                                                                                                                                                                     /label= Signal_peptide
/note= "amino acids 1-7 are PCR primer-encoded
differ from the native sequence"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antibody
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                                                               TNV14 heavy chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           necrosis factor anti
es, atherosclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human IgG1, kappa isotype. They showed relate the human germline DP-46 heavy chain sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'n
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                infarction,
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                                                                                                                                                                                                                                                                  Mature protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Heavner
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                                                                                                                                                                                              Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        English
146
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2000US-0236826P.
2001US-00920137.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cell anemia,
                                                                                                                                                                                                                                                                                                                                                         CDR2
                                                                                                                                                                                                                                                                                                              CDR1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001WO-US024785
                                                                                                                                                                                                                                                                                                                                                                  86. .117
/label= FR3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   myocardial
protein;
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                                                                                                                                                                                                                                                                                                                                                                                         135
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                                                                                                                                                                                                                                                                                                                                            09. .85
/label=
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/label=
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/label=
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/label=
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/label=
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P
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                                                                                                                                                                                                                                                                                                                                                                                          118
                                                                                                                                                                                                                                                                                                                                                                                                               56
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                                           (first
                                                                                                                                                                                                                                                       20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002-217194/27
standard;
                                                                Human recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    angina pectoris,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB; ABL53508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treating sickle
                                                                                      Tumour necrosis complementarity
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                                                                                                                               antianginal; coprotozoacide; human; diagnos
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                                                                                                                                                                            sapiens.
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01-AUG-2001;
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AAM51168
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may

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other

(mAb)

invention provides isolated human, primate, rodent, mammalian, chimeric, humanised and/or complementarity determining region (CDR)-grafted antiTNF antibodies, immunoglobulins, cleavage products and other specified portions and variants, as well as anti-TNF antibody compositions, complementary nucleic acids, vectors, host cells, compositions, formulations, devices, transgenic animals, transgenic compositions, formulations devices, transgenic animals, transgenic plants, and methods of making and using them. The anti-TNF antibody comprises at least a portion of an immunoglobulin molecule, especially the heavy chain and/or light chain variable regions given in the present sequence and in AAM51165, or either all of the CDRs of the heavy chain (see AAM51165, or either all of the CDRs of the heavy chain (see AAM5118-60) or all of the CDRs of the light chain (see AAM51161-63). The antibodies may inhibit TNF-induced cell adhesion molecules, in a mouse model. They are useful for diagnosing or treating a TNF related condition in a cell, tissue, organ or animal (claimed) such as theumatoid arthritis, gastric ulcer, asthma, allergic rhinitis, Crohn's pathology, sickle cell anaemia, diabetes, a cardiovascular disease such as arteriosclerosis, atherosclerosis, restenosis, angina pectoris or myocardial infarction, an infectious disease in a cell such as bacterial, viral, and fungal infections, pneumonia, leprosy and malaria, a malignant disease such as leukaemia, chronic myelocytic leukaemia, Burkitt's lymphoma and multiple myeloma, or a neurological disease such as multiple sclerosis, parkinsor's disease, spinal ataxia, Alzheimer's disease and i, 119 receptor; 9 79 Nile virus; WNV. QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY 20 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLEWVAIILYDGSSKKY Human heterodimeric antibody heavy chain variable region SEQ ID NO:98 human heterodimeric antibody; human; antibody; binding affinity; protective antigen; Bacillus anthracis; anthrax infection; cell recepedema factor; lethal factor; virucide; antibacterial; immunotherapy; anti-toxin; anti-infective; anthrax; botulinum; smallpox; Venezuelan equine encephalomyelitis virus; VEEV; West Nile virus; WNN Gaps ; 1 146; Length Indels 5 5; DB Score 615.5; DB Pred. No. 3e-45; ADD28320 standard; protein; 135 AA э ; 2002US-0356086P. 2002US-0376408P. 2002US-0414053P. 2002US-0428807P. 92.1%; ilarity 92.9%; Conservative 2003WO-US004206 entry) Creutzfeldt-Jakob disease 146 TTVTVSS 126 (first l Similarity 118; Conser TTVTVSS Sequence 146 AA; WO2003076568-A2 11-FEB-2002; 29-APR-2002; 27-SEP-2002; 25-NOV-2002; 11-FEB-2003; Homo sapiens 18-SEP-2003 5-JAN-2004 Synthetic ADD28320; 140 120 Query Match Best Local Н Matches ADD28320 RESULT 88888888888888888888888888888888888 셤 셤 В ਨੇ 8 ઠે

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26-MAY-2004;
27-SEP-2002;
25-NOV-2002;
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Best Local S
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                                                                   useful as
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tis or West
                                                                                                                              The present invention describes a human heterodimeric antibody (I) (fragment) having a binding affinity of at least 1x10-8 M to the protective antigen of Bacillus anthracis or a molecule involved in anthrax infection that blocks binding of the antigen or molecule to cell receptors, edema factor and lethal factor. (I) has virucide and antibacterial activities, and can be used in immunotherapy. The antibodies (I) are useful as anti-toxins or anti-infectives with respect to infective agents, such as anthrax, botulinum, smallpox, Venezuelan equine encephalomyelitis virus (VEEV), or West Nile virus (WNV). The present sequence represents a human heterodimeric antibody heavy chain variable region amino acid sequence, which is used in the exemplification of the present invention.
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                                                                              agents
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                                                                   or their antibody respect to infec
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1.7e-42;
                                                                                                                                                                                                                                                                                          Score 583; DB Pred. No. 1.7e-4; Mismatches
                                                                   New human heterodimeric antibodies or their anti-toxins or anti-infectives with respect
                                                                                                                     Claim 11; SEQ ID NO 98; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           equine encephalomyelitis
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2002US-0376408P
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86.7%;
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          (ALEX-) ALEXION PHARM INC
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                              wild MA;
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29-APR-2002;
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                                                                                                  Nile virus.
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The present invention describes a human heterodimeric antibody (I) (fragment) having a binding affinity of at least 1x10-8 M to the protective antigen of Bacillus anthracis or a molecule involved in anthrax infection that blocks binding of the antigen or molecule to cell receptors, edema factor and lethal factor. (I) has virucide and antibacterial activities, and can be used in immunotherapy. The antibodies (I) are useful as anti-toxins or anti-infectives with respect to infective agents, such as anthrax, botulinum, smallpox, Venezuelan equine encephalomyelitis virus (VEEV), or West Nile virus (WNV). The present sequence represents a human heterodimeric antibody heavy chain variable region amino acid sequence, which is used in the exemplification of the present invention.
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2002US-0414053P
2002US-0428807P
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ilarity 86.7%;
Conservative
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antibacterial; antibody
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                                                                                                               Wild MA;
                                                                                                                                                           2003-722327/68
                                                                                                                                                                                                                                                                                                                       Claim 6; SEQ ID NO
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acid sequence

amino

chain

T3F2 heavy

Human antibody

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comprises administering an animal infected with Bacillus anthracis comprises administering an antibody or antibody fragment that binds to a molecule involved in anthrax infection and which has the ability to block the binding of the molecule to at least one of a cell receptor. A claimed method for the binding of the molecule to at least one of a cell receptor. A claimed method for determining exposure to B. anthracis comprises assaying a sample for the presence of a molecule selected from cell receptors, PA63, PA63 heptamer, PA83, edema factor or lethal factor with an antibody that has binding affinity for the molecule, where the presence of a disease associated with B. antibody correlates with the presence of a disease associated with B. antibody correlates with the spresence of a leaveted levels of the accondary cort the antibody, where the presence of B. anthracis in a subject. In all cases, the antibody (full-length or functional fragment) may comprise a heavy chain variable region selected from a group or sequences ADV86812, a light chain lambda region selected from a group or sequences ADV86823 and a light chain lambda region selected from a group or sequences ADV86836.ADV86847. Diagnostic kits are provided. A claimed vaccine comprises a multimer of anthrax toxin PA63. The methods and compositions of the present invention are also useful for producing anticompositions of the present invention are also useful for producing anticompositions or anti-infectives to infective agents such as anthrax, botulinum, smallpox, venezulan equine encephalomylain variable region of the heavy chain variable region of the heavy chain variable region are compositions of the heavy chain variable region are cellulated from a group of present sequence is that of the heavy chain variable region are cellulated from thrax provided. The present sequence is tha
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nding to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        against anthrax. The libraries were panned against PA83 and PA63, and sequence analysis was performed on positive responders. Neutralization anthrax toxin activity by purified Fabs was demonstrated.
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bin
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proteins PA63 and PA83. Phage libraries were developed from mR
from blood and bone marrow samples of donors who had been vacc
                                                                                                             Nolan MJ
                                                                                                                                                                                                                     Treating an animal with anthrax infection by administering the protective antigen of Bacillus anthracis, and blocking cell receptors, edema factor or lethal factor.
                                                                                                             H
                                                                                                              Maruyama
                                                                                                           Wild MA,
                                                                                                                                                                                                                                                                                                                                    Claim 7; SEQ ID NO 14; 87pp; English
    2003US-00452593
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                                                        (ALEX-) ALEXION PHARM INC.
                                                                                                             Prederickson
                                                                                                                                                                  2005-057715/06
02-JUN-2003;
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GMDVWGQ IIII I I ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARDRGIAAGGN--YYYYGMDVWGO **QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY** 5 Length 135; Indels 11; , DB 9; 1.7e-42; ed. No. 1.76 Mismatches Score 583; Pred 4; 87.3**%**; 86.7**%**; Conservative Local Similarity nes 111; Conser Ä Sequence 135 61 Н Query Match Matches <del>Q</del> a Š ઠ

ADA89238 standard; protein; 124 AA. (first entry) 126 GTTVTVSS 20-NOV-2003 ADA89238; 119 RESULT 9 ADA89238 XXXEX П 요 8

domain for disorder. immunoglobulin; Ig; heavy chain variable domain; light chain variable domain; major histocompatibility complex; MHC; gp100; MUC1; TAX; hTERT; cytostatic; gene therapy; cancerous disorder; New protein comprising an immunoglobulin heavy chain variable (VH) and an immunoglobulin light chain variable (VL) domain, useful for preparing a composition for treating or preventing a cancerous disc TECHNION RES & DEV FOUND LTD. 20-FEB-2002; 2002US-0358994P 2003WO-US005128 Reiter Y; 2003-663847/62. CORP N-PSDB; ADA89237 Hoogenboom HRJM, WO2003070752-A2 20-FEB-2003; (DYAX-) DYAX Homo sapiens 28-AUG-2003 Synthetic (TECR ) cancer 

Disclosure; Fig 19B; 224pp; English.

The present invention describes a protein comprising an immunoglobulin (Ig) heavy chain variable (VH) domain and an Ig light chain variable (VL) domain. The protein binds a complex comprising a major histocompatibility complex (MHC) and a peptide, does not substantially bind the MHC in the absence of the bound peptide, and does not substantially bind the Peptide in the absence of the MHC. The peptide is a peptide fragment of gpl00, MUC1, TAX or hTERT. Also described: (1) a pharmaceutical composition comprising the novel protein and a carrier; (2) a cytocoxic T cell comprising one or more nucleic acids for expressing the Ig that binds a complex having an MHC and a peptide, does not substantially bind the mace of the bound peptide, and does not substantially bind the comprising a first segment that encodes the Ig variable domain; (4) a host cell comprising heterologous nucleic acid sequences that encodes the Ig variable domain; (5) a transgenic animal whose genome includes heterologous nucleic acid sequences that encode the protein; (6) identifying the protein that specifically binds the MHC-peptide complex; (7) expressing an antigen-binding protein; (8) ablating or killing a target cell that displays a peptide on a surface MHC molecule; (9) treating or preventing an antigen-binding protein; (8) ablating or killing a target cell that displays and suspect of the invention has cytostatic activity, and can be used in gene therapy. The protein is useful for preparing a composition for treating or preventing a cancerous disorder. The present peptide complex where the peptide component in as peptide fragment ine present

of

7 Length 124; Indels 9; 9 Score 582; DB 6 Pred. No. 2e-42; Mismatches 3; 87.1%; 88.9%; Conservative Similarity 112; ery Match Matches 8

Sequence 124 AA;

118

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9 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT -61

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Best Local S
Matches 111
            121
                                                                     AAW80815
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(RAUM/)
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ID AEC16
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                                                RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a method of identifying binding site domains

(BSD) that retain the capacity of binding to a predetermined epitope when

positioned C-terminal of at least one further domain in a recombinant bi-

or multivalent polypeptide. The method comprises (a) testing a panel of

BSD displayed on the surface of a biological display system as part of a

fusion protein for binding to a predetermined epitope, where the fusion

protein comprises an additional domain positioned N-terminal of the BSD

and an amino acid sequence that mediates anchoring of the fusion protein

to the surface of the display system; and (b) identifying a BSD that

binds to the predetermined epitope. The method is useful to identify bi-

c or multivalent polypeptides that comprise antibody binding sites capable

or multivalent polypeptides that comprise antibody binding sites capable

c of efficiently binding to the corresponding antigen. The polypeptides or

antibodies identified by the method are useful therapeutically and

diagnostically, for e.g. cancer and autoimmune diseases. ScFv-antibody

fragments that bind independently of their position within bifunctional

single-chain fusion proteins can be isolated from combinatorial antibody

libraries using the new in vitro method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ή,
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                                                                                                                                                                            cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                           domains
                                                                                                                                                                         fusion protein; therapeutic single-chain Fv.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 127;
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                                                                                                                                                                                                                                                                                                                                                                                                          identification of binding site
                                                                                                                                                                                                                                                                                                                                                   Lutterbuese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 581.5; DB 2;
Pred. No. 2.2e-42;
; Mismatches 10;
                                                                                                                                                                                                                                                                                                                                                  F,
                                                                                                                                                    Human D4.5 heavy chain variable region
                                                                                                                                                                                                                                                                                                                                                  Zett1
                                                                                                                                                                                                                                                                                                                                                                                                                                             3.1; 152pp; English.
                                                                                                                                                                         Binding site domain; BSD; epitope; autoimmune disease; scFv-antibody;
                                                                                standard; protein; 127
                                                                                                                                                                                                                                                                                                                                                  Χ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5;
                                                                                                                                                                                                                                                                                                     97EP-00120096.
                                                                                                                                                                                                                                                                                                                                                  Borschert
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ilarity 87.4%;
Conservative
                                                                                                                                                                                                                                                                               98WO-EP007313
                                                                                                                                                                                                                                                                                                                                                                                                                      epitope
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                                                                                                                                                                                                                                                                                                                                                                                                           Phage display system for capacity to bind an epitc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         120 TTVTVSS 126
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                  Raum T,
                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB; AAX77236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 127 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig
TVTVSS
                        TVTVSS
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                                                                                                                                                                                                            sapiens
                                                                                                                                                                                                                                   WO9925818-A1
                                                                                                                                                                                                                                                                               16-NOV-1998;
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                                                                                                                             04-AUG-1999
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                                                                                                       AAY17954;
 121
                        119
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This is the amino acid sequence of the human D4.5 heavy chain variable region, used in the method of the invention. for providing receptors that can be used for targeting antigens in humans without being immunogenic themselves. Such receptors can be used for treating diseases such as tumours or auto-immune diseases, graft rejection after transplantation, infectious diseases by targeting cellular receptors as well as allergic, inflammatory, endocrine and degenerative diseases by targeting key molecules involved in the pathological process
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      duction of anti-human antigen receptors - by selecting a combination functionally rearranged VH and VL immunoglobulin chains expressed from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARDRGIAAGGN-YYYYGMDVWGQG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tumour;
disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                     Amino acid sequence of human D4.5 heavy chain variable region.
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                                                                                                                                                                                                                                                                                                Human; D4.5 heavy chain variable region; receptor; antigen; auto-immune disease; graft rejection; allergy; inflammatory endocrine disease; degenerative disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 581.5; DB 2;
Pred. No. 2.4e-42;
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                                                                                                                standard; protein; 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2
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87.4%;
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127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Raum T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TTVTVSS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Д
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                                                                                                                                                                                                                                                                                                                                                                                          sapiens
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                                                                                                                                                                                                          16-FEB-1999
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HD70; single-chain variable fragment; scFv; 17-1A antigen; human; EpCAM; epithelial cell adhesion molecule; inflammatory cytokine; GM-CSF; granulocyte/macrophage colony stimulating factor; heterominibody; CH1-domain; multifunctional compound; heavy chain constant domain; immunoglobulin; cytostatic; immunostimulatory; antileukaemia; diagnosis; antiproliferative; prevention; treatment; malignant; haematopoietic cell lymphoma; leukaemia; solid tumour; carcinoma; melanoma; sarcoma.

g

AAY44994 standard; protein;

AAY44994

entry)

(first

23-MAY-2000

AAY44994;

HD70scFv-CH1-GM-CSF chain.

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The invention relates to a method of treating tumorous disease by administering a human immunoglobulin specifically binding to the human epithelial cell adhesion molecule (BpCAM; also termed as 17-1A antigen, KSA, EGP40, GA733-2, ks 1-4 and esa) antigen. The immunoglobulin exhibits a serum half-life of at least 15 days and is administered no more frequently than once every week. The method of the present invention and composition comprising anti-EpCAM are useful for treating tumorous disease, such as breast cancer, epithelial cancer, hepatocellular cancer, eromach cancer, negative cancer, head and neck cancer, stin cancer, colon cancer, cancer, and kidney cancer, lung cancer, skin cancer, a cancer of the small intestine, liver cancer, pancreas cancer, gall bladder cancer, a cancer of the bile duct, esophagus cancer, a cancer of the salivary cancer of the thyroid gland. The present sequence is the human anti-EpCAM immunoglobulin heavy chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       adhesion
                                                                                                                                                                                               ;
bulin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     liver,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cell
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                                                                                                                   Antibody therapy; tumor; cancer; cytostatic; neoplasm; pancreas ovary tumor; thyroid tumor; stomach tumor; breast tumor; epithelial cancer; hepatocellular carcinoma; cholangiocellular head & neck tumor; skin tumor; melanoma; uterine cancer; uterine cervix tumor; renal tumor; lung tumor; intestine tumor liver tumor; gallbladder carcinoma; esophagus tumor; immunoglol antibody; epithelial cell adhesion molecule; heavy chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating tumorous disease, such as breast, colon, prostate, ovarian, cervical and lung cancer, by administering a human immunoglobulin specifically binding to the human epithelial molecule (EpCAM) antigen.
                                                                                    Human anti-EpCAM immunoglobulin heavy chain, SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Quadt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 1; 23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prang N,
                                                                                                                                                                                                                                                                                                                                                                        2004US-00778915
                                                                                                                                                                                                                                                                                                                                                                                                           13-FEB-2004; 2004US-00778915
                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Locher M,
                                                                                                                                                                                                                                                                                                                                                                                                                                             (MICR-) MICROMET AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2005-590351/60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 457 AA;
                                                                                                                                                                                                                                                                                                   US2005180979-A1
                                                                                                                                                                                                                                                                 sapiens
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                                                  20-OCT-2005
                                                                                                                                                                                                                                                                                                                                      18-AUG-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 12;
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2;
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                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                          Sequence 523 AA;
                                                                                                                                                                                                                                                           111;
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                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                       ADSVKGRPTISRDNSKNTLYLQMNSLRAEDTAVYYCAKDMGWGSGWRPYYYYGMDVWGQG
                                                                                                                                                                                                                                 OVOLVESGGGVVOPGRSLRLSCAASGFIPSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                         1;
                                                                                                                                                                                                                457;
                                                                                                                                                                                                                Length
                                                                                                                                                                                                                         Indels
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3.1e-42;
1es 10;
                                                                                                                                                                                                               Score 581.5; I
Pred. No. 8.1e-
5; Mismatches
                                                                                                                                                                                                                         5;
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illarity 87.4%;
Conservative
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Similarity

Query Match Best Local S

111;

Matches

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TTVTVSS

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produceable in a mammalian host cell as a secretable and fully functional certains produceable in a mammalian host cell as a secretable and fully functional heterodimer of two polypeptide chains, where one of the polypeptide chains comprises, a CH1-domain (constant domain of an immunoglobulin and the other chain comprises CL-domain (constant domain of an immunoglobulin light chain). The polypeptide chains further comprise, can immunoglobulin light chain). The polypeptide chains further comprise, different receptor or ligand functions, where further at least two of the different (poly)peptides lack an intrinsic affinity for one another and are linked via the constant domains. The heterominibodies have cytostatic, immunostimulatory, antileukaemia and antiproliferative activities. These compounds can be used for diagnosing, preventing and treating malignant cell growth related to malignancies of haematopoletic cells e.g. lymphomas and leukaemias, or to solid tumours e.g. carcinomas, melanomas and sarcomas. The present sequence is the left chain of a heterominibody comprising HD70 single-chain PV (scFv) fragment N-terminally linked to human CH1 domain which bears at its C-terminus the human inflammatory cytokine granulocyte/macrophage colony stimulating factor (GM-CSF), plus a hexahistidine sequence for ease of purification. CHD70 scFv specifically recognises the human epithelial cell adhesion colons.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          201
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Pred. No. 9.3e-42;
166pp; English
    55A;
  Example 10; Fig
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treating

and/or

New multifunctional compounds useful for preventing an malignant cell growth and for detection and diagnosis.

Zettl

Borschert K,

Baeuerle PA,

H

Dreier

<u>ц</u>

Kufer

8kin,

WPI; 2000-195265/17 N-PSDB; AAZ50587.

BIOMEDIZINISCHE FORSCHUNG

GES

(MICR-) MICROMET

99WO-EP005416

28-JUL-1999;

WO200006605-A2

10-FEB-2000.

Homo sapiens.

98EP-00114082

28-JUL-1998;

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The patent discloses heterominibodies which are multifunctional compounds produceable in a mammalian host cell as a secretable and fully functional broaduceable in a mammalian host cell as a secretable and fully functional heterodimer of two polypeptide chains, where one of the polypeptide chains comprises, a CH1-domain (constant domain of an immunoglobulin and the other chain comprises CL-domain (constant domain of an immunoglobulin light chain). The polypeptide chains further comprise, fused to the constant domains at least two (poly)peptides having different (poly)peptides lack an intrinsic affinity for one another and are linked via the constant domains. The heterominibodies have cytostatic, immunostimulatory, antileukaemia and antiproliferative activities. These compounds can be used for diagnosing, preventing and treating malignant cell growth related to malignancies of haematopoietic cells e.g. lymphomas and leukaemias, or to solid tumours e.g. carcinomas, melanomas and sarcomas. The present sequence is the right chain of a heterominibody comprising HD70 single-chain Fv (scFv) fragment N-
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functional
119
                                                            261
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1ymphoma
MDVWGQG
                             HD70; single-chain Fv fragment; scFv; antibody; 17-1A antigen; human; EpCAM; epithelial cell adhesion molecule; inflammatory cytokine; IL-2; interleukin-2; Ck-domain; kappa light chain constant domain; heterominibody; multifunctional compound; immunoglobulin; cytostatic; immunostimulatory; antileukaemia; diagnosis; prevention; antiproliferative; treatment; malignant; haematopoietic cell; lymphom; leukaemia; solid tumour; carcinoma; melanoma; sarcoma.
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                                                                                                 61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT
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                                       OVOLVESGGGVVOPGRSLRLSCAASGFIFSSYAMHWVROAPGNGLEWVAFMSYDGSNKKY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New human monoclonal antibody that binds to platelet-derived factor-D (PDGF-D), useful for treating chronic and recurrent diseases, such as inflammation, autoimmunity and cancer.
                                                                                                                                                                                                                                                                                                                  immunomodulator; cytostatic; gene therapy
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Search completed: December 28, 2005, 13:58:06 Job time : 149.538 secs

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GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd
                   Copyright
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sw model using - protein search, OM protein ; Search time 25.8462 Seconds (without alignments) 469.057 Million cell updates/sec 13:46:34 2005, 28, December Run on:

US-09-920-137F-7 668 score: Perfect

.....NYYYYGMDVWGQGTTVTVSS **QVQLVESGGGVVQPGRSLRL...** Н Sequence:

126

Gapext 0.5 BLOSUM62 Gapop 10.0 , Scoring table:

residues 283416 seqs, 96216763 Searched:

283416 of hits satisfying chosen parameters: Total number

Minimum DB seq Maximum DB seq

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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

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pirl: \*
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pir3: \* 1 0 m 4

Pred. No. is the number of results predicted by chance to have score greater than or equal to the score of the result being prand is derived by analysis of the total score distribution.

#### SUMMARIES

	Description	g heavy chain V	g heavy chain V	g heavy chain -	avy	g heavy chain -	g heavy chain	g variable regio	g heavy chain V-	g heavy chain V	g heavy chain pr	g heavy chain V	g heavy chain -	g heavy chain -	g heavy chain pr	g heavy chain	g heavy chain V	g heavy chain V	g heavy chain pr	g mu chain - hum	g heavy chain V	g heavy chain V	g heavy chain V	g heavy chain	g heavy chain V	g heavy chain V	g heavy chain V			
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## ALIGNMENTS

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RESULT 1
E36005
Ig heavy chain V region (M72) - human
C;Species: Homo sapiens (man)
C;Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 16-Dec-:
C;Accession: E36005
R;Schroeder Jr. H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy cha. A;Reference number: A36005; MUID:90349571; PMID:2117273
A;Actus: preliminary
A;Molecule type: mRNA
A;Residues: 1-122 <SCH>
A;Cross-references: UNIPARC:UPI0000176C30; GB:M34030
C;Genetics:
A;Gene: GDB:IGH@Y; IGHDY1
A;Cross-references: GDB:118731; OMIM:146910
A;Map position: 14q32.33-14q32.33
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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122 TVTVSS 117 RESULT S48797 Ig heav 셤

Ig heavy chain V region (anti-Sm, VH3/Dxp4/JH6) - human (fragment)
C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 13-Jan-1995 #sequence\_revision 13-Sep-1998 #text\_change 23-Ju1-1999
C; Accession: S48797; S26893
R; Mahmoudi, M.; Edwards, J.; Cairns, E.; Bell, D.
Submitted to the EMBL Data Library, October 1994
A; Description: Molecular characterization of natural human anti-Sm autoantibodies A; Reference number: S48797
A; Accession: S48797
A; Molecule type: mRNA

gene

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M.J.D.; Vossen, J.M.; Schuurman
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R;Schroeder Jr., H.W.; Wang, J.Y.

Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy char.
A;Accession: F36005
A;Accession: F36005
A;Status: preliminary
A;Accession: F36005
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-119 <SCH>
A;Cross-references: UNIPROT:QBWUK1; UNIPARC:UPI0000176C32; GB:M34026
C;Genetics:
A;Gene: GDB:IGHØ; IGHDY1
A;Cross-references: GDB:118731; OMIM:146910
A;Genetics:
C;Superfamily: immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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Pred. No. 2.6e-41;
2; Mismatches 10;
                                                                                                                                                                                                                                                         539.5; DB No. 1.5e-41
                                                                                                                                                                                                                                                                                       Mismatches
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Pred.
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84.1%;
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Best Local Similarity 84.1%;
Matches 106; Conservative
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Best Local Similarity
Matches 106; Conser
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                                                                                                                                         PIDN: CAA78220.1;
                                                                   groups
                      PIDN:CAA86512
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   A; Residues: 1-128 < MAH>
A; Cross-references: UNIPARC: UPI0000116700; EMBL: Z46379; NID: 9587147; PI)
R; Tomlinson, I.M.; Walter, G.; Marks, J.D.; Llewelyn, M.B.; Winter, G.
J. Mol. Biol. 227, 776-798, 1992
A; Title: The repertoire of human germline V(H) sequences reveals about A; Reference number: S26885; MUID: 93021117; PMID: 1404388
A; Residues: 1-98 < TOM>
A; Residues: 1-98 < TOM>
A; Cross-references: UNIPARC: UPI0000038183; EMBL: Z12350; NID: 932922; PID
C; Superfamily: immunoglobulin V region; immunoglobulin
C; Keywords: heterotetramer; immunoglobulin
F; 15-98/Domain: immunoglobulin homology < IMM>
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C; Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_c; Accession: 838493
R; Marks, J.D.; Ouwehand, W.H.; Bye, J.M.; Finnern, R.; Gorsubmitted to the EMBL Data Library, June 1993
A; Description: Human antibody fragments specific for human A; Reference number: 838488
A; Accession: 838493
A; Accession: 838493
A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-123 < MAR>
A; Cross-references: UNIPARC: UPI000011654F; EMBL: Z23036; NI
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C;Superfamily: immunoglobulin V region; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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Pred. No. 1.1e-43;
                                                                                                                                                                                                               ced. No. 1.1e-44;
Mismatches 11
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                                                                                                                                                                                                               Score
Pred.
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nilarity 87.5%;
Conservative
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nilarity 87.3%;
Conservative
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F36005
Ig heavy chain V region (M49) - C;Species: Homo sapiens (man) C;Date: 21-Dec-1990 #sequence_r
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hes 110; Conser
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S38493
Ig heavy chain -
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G; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 21-Jan-1994 #sequence_revision 18-Nov-1994 #text_change 23-Jul-1999
C; Date: 21-Jan-1994 #sequence_revision 18-Nov-1994 #text_change 23-Jul-1999
C; Accession: A49028
R; Timmers, E.; Kenter, M.; Thompson, A.; Kraakman, M.E.; Berman, J.E.; Alt, F.W.; Schuur, Eur. J. Immunol. 21, 2355-2363, 1991
A; Title: Diversity of immunoglobulin heavy chain gene segment rearrangement in B lymphob A; Reference number: A49028; MUID: 92008140; PMID: 1915549
A; Reference number: A49028
A; Retus: preliminary
A; Residues: 1-133 <TIM>
A; Residues: 1-133 <TIM
A; Residues: 1-133 <TIM
A; Residues: 1-133 <TIM
A; Residues: 1-13028 <TIM
A; Residues: 1-13028 <TIM
A; Residues: 1-13028 <TIM
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C;Species: Homo sapiens (man)
C;Date: 27-Jan-1995 #sequence_revision 27-Jan-1995 #text_change 20-Jun-2(C;Date: 27-Jan-1995 #sequence_revision 27-Jan-1995 #text_change 20-Jun-2(C;Accession: S46390
R;Figini, M.; Marks, J.D.; Winter, G.; Griffiths, A.D.
J. Mol. Biol. 239, 68-78, 1994
A;Title: In vitro assembly of repertoires of antibody chains on the surfa
A;Reference number: S46390; MUID:94254092; PMID:8196048
A;Reference number: S46390
A;Residues: preliminary
A;Molecule type: DNA
A;Residues: 1-114 <FIG>
A;Cross-references: UNIPARC:UPI000011663F; EMBL:Z31686; NID:g509782; PIDP
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
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. No. 6.9e-41;
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Pred. No. 7.3e-41;
2; Mismatches 12;
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illarity 83.3%;
Conservative
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107; Conservative
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Best Local Similarity
Matches 105; Conser
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            lg neavy chain v region (m/4) - nummain
C;Species: Home sapiens (man)
C;Species: 12-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 31-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 31-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 31-Dec-1990 #schroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin heavy cha A;Reference number: A36005; MUID:90349571; PMID:2117273
A;Reference number: A36005; MUID:30349571; PMID:2117273
A;Reference number: A36005; MUID:30349571; PMID:2117273
A;Refe
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C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 16-Feb-1996 #sequence_revision 16-Feb-1996 #text_change 23-Jul-C; Accession: I37780; S25474
R; Demaison, C.; Chastagner, P.; Theze, J.; Zouali, M.
R; Demaison, C.; Chastagner, P.; Theze, J.; Zouali, M.
Proc. Natl. Acad. Sci. U.S.A. 91, 514-518, 1994
A; Title: Somatic diversification in the heavy chain variable region general approachment: A36876; MUID:94119917; PMID:8290556
A; Accession: I37780
A; Accession: I37780
A; Accession: I37780
A; Accession: I37780
A; Molecule type: mRNA
A; Residues: 1-147 < RES>
A; Cross-references: UNIPARC:UPI00001160E7; EMBL:X67943; NID:g33578; PID:C; Superfamily: immunoglobulin variable rimmunoglobulin homology < IMM>
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                         Igheavy chain V region - human
C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 03-Mar-1994 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C; Accession: S31603
R; Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
submitted to the EMBL Data Library, June 1992
A; Description: Mechanisms that generate human immunoglobulin diversity operate from A; Reference number: S31585
A; Accession: S31603
A; Accession: S31603
A; Accession: S31603
A; Residues: 1-132 <CUI>A; Residues: 1-132 <CUI>A; Cross-references: UNIPARC:UPI0000116455; EMBL:Z14168; NID:g30999; PIDN:CAA78537.1
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 30-113/Domain: immunoglobulin homology <IMM>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ig heavy chain V region (VH3DJH4) - human

C; Species: Homo sapiens (man)

C; Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change 20-Jun-C; Accession: S1966

R; Marks, J.D.; Hoogenboom, H.R.; Bonnert, T.P.; McCafferty, J.; Griffit J. Mol. Biol. 222, 581-597, 1991

A; Title: By-passing immunization. Human antibodies from V-gene librarie A; Reference number: S19663; MUID: 92085276; PMID: 1748994

A; Accession: S19666

A; Molecule type: mRNA

A; Residues: 1-121 < MAR>
A; Residues: 1-121 < MAR>
A; Cross-references: UNIPARC: UPI0000115FE5; EMBL: X61646; NID: g37688; PIC C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 15-98/Domain: immunoglobulin homology < IMM>
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Pred. No. 1.2e-40;
5; Mismatches 11;
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                                                                                                                                                                                                                                                                                                      DB
                                                                                                                                                                                                                                                                                                    Score 530.5; I
Pred. No. 1.1e-
4; Mismatches
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Local Similarity 82.5%;
nes 104; Conservative
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Best Local Similarity 82.5%;
Matches 104; Conservative
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Best Local S:
Matches 104
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C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-; C; Accession: S31679
R; Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C submitted to the EMBL Data Library, June 1992
A; Description: Mechanisms that generate human immunoglobulin diversity of A; Reference number: S31585
A; Reference number: S31585
A; Accession: S31679
A; Accession: S1679
A; Residues: 1-134 «CUI»
A; Residues: 1-134 «CUI»
A; Residues: 1-134 «CUI»
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
C; Keywords: heterotetramer; immunoglobulin
F; 34-117/Domain: immunoglobulin homology <IMM>
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                                                                                                                    immunoglobulin diversity
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C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 24-Jul-1998 #sequence_revision 24-Jul-1998 #text_chang
C; Accession: S70442
R; Cuisinier, A.M.; Fumoux, F.; Fougereau, M.; Tonnelle, C.
Mol. Immunol. 29, 1363-1373, 1992
A; Title: IgM kappa/lambda EBV human B cell clone: an early ste
A; Reference number: S70442; MUID: 93024508; PMID: 1383695
A; Accession: S70442
A; Accession: S70442
A; Molecule type: mRNA
A; Molecule type: mRNA
A; Residues: 1-140 <CUI>A; Residues: 1-140 <CUI>A; Cross-references: UNIPROT: Q8WUK1; UNIPARC: UPI0000176EB7
C; Superfamily: immunoglobulin homology
F; 34-117/Domain: immunoglobulin homology
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Pred. No. 1.7e-40;
2; Mismatches 6
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Pred. No. 2e-40;
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S46392
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of

Ig heavy chain V region (VH-28)

12

RESULT

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PIDN: CAA83493.1; PI
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C; Species: Homo sapiens (man)
C; Date: 27-Jan-1995 #sequence_revision 27-Jan-1995 #text_change 20-Jun-;
C; Accession: 846392
R; Figini, M.; Marks, J.D.; Winter, G.; Griffiths, A.D.
J. Mol. Biol. 239, 68-78, 1994
A; Title: In vitro assembly of repertoires of antibody chains on the sur. A; Reference number: 846390; MUID:94254092; PMID:8196048
A; Accession: 846392
A; Accession: 846392
A; Accession: 846392
A; Molecule type: DNA
A; Molecule type: DNA
A; Residues: 1-114 <FIG>A; Residues: 1-114 <FIG>A; Cross-references: UNIPARC:UPI00001137D6; EMBL:231688; NID:9499306; PIC; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 15-98/Domain: immunoglobulin homology <IMM>
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C; Species: Homo sapiens (man)
C; Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-C; Accession: S31117
R; Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vo Eur. J. Immunol. 22, 247-251, 1992
A; Title: Restricted utilization of germ-line V(H)3 genes and short A; Reference number: S31104; MUID:92111633; PMID:1730252
A; Reference number: S31117
A; Reference number: S31117
A; Residues: 1-122 <RAA>
A; Residues: 1-122 <RAA>
A; Cross-references: UNIPARC:UPI0000176C8D; EMBL:X62967
A; Note: the nucleotide sequence was submitted to the EMBL Data Libr C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 15-98/Domain: immunoglobulin homology <IMM>
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Pred. No. 3.9e-40;
H; Mismatches 13
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Pred. No. 3.6e-40,
1; Mismatches
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Best Local Similarity 82.5%;
Matches 104; Conservative
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Best Local Similarity 83.3%;
Matches 105; Conservative
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#### Protein Sequence Searches - February 2005

All of the sequence databases on ABSS have recently been updated.

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- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt record may not contain the previous temporary accession number.

  accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.

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Compugen Ltd
version:
GenCore (c) 1993
        Copyright
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model 3 using protein search, OM protein

; Search time 159.923 Seconds (without alignments) 555.871 Million cell updates/sec 13:44:24 2005, 28, December . : Run

US-09-920-137F-7

899 score: Perfect

.....NYYYYGMDVWGQGTTVTVSS QVQLVESGGGVVQPGRSLRL.. Sequence:

126

BLOSUM62 Gapop 10.0 0 table Scoring

Gapext 0.5

residues 705528306 2166443 seqs, Searched

2166443 of hits satisfying chosen parameters Total number

2000000000 length: length: seq seq 88 Minimum Maximum **%** 

Maximum Match 100% Listing first 45 summaries Post-processing: Minimum Match Maximum Match

UniProt Database

iProt\_05.80:\*
uniprot\_sprot:\*
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printed, Pred. No. is the number of results predicted by chance to have score greater than or equal to the score of the result being prand is derived by analysis of the total score distribution.

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#### ALIGNMENTS

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TISSUE=Primary B-Cells;

MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Strausher R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Stapleton M., Soares M.B., Peters G.J., Abramson R.D., Mullahy S.J.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Rosark S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Rahey J., Helton B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human
                                                                                                                                                                                   Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ion of conserved immunoglobulin heavy chain during human fetal life."; U.S.A. 87:6146-6150(1990).
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Sci. U.S.A. 99:16899-16903(2002)
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PubMed=1383695; DOI=10.1016/0161-5890(92)90173-U;
Cuisinier A.M., Fumoux F., Fougereau M., Tonnelle
                                         613 AA
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PubMed=2117273;
Schroeder H.W. Jr, Wang J.Y.;
"Preferential utilization of
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26,
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                                      QBWUK1 HUMAN PRELIMINARY;
QBWUK1;
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01-MAR-2002 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
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TISSUE=Primary B-Cells;
NIH MGC Project;
Submitted (DEC-2001) to
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Proc. Natl. Acad. Sci.
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Name=IGHM;
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Millson J.L., Karr N.S., Oppliger I.R., Mannik M., Sasso E.H.;

"The structural basis of germline-encoded VH3 immunoglobulin bindi
to staphylococcal protein A.";

J. Exp. Med. 178:331-336(1993).

R EMBL; BC020240; AAH20240.1; -; mRNA.

R PIR; P36005; P36005.

R PIR; PH1642; PH1642.

R PIR; PH1642; PH1642.

R PIR; PH1645; PH1645.

R PIR; PH1645; PH1646.

R PIR; S31119; S31119.

R PIR; S31119; S31119.

R PIR; S70442; S70442.

R SSR; Q8WUKL; 20-242.

R ENSEMb; ENSG00000130076; Homo sapiens.
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Dard J., Galili N., Link M., Stites D., Sklar J.;
"Continuing rearrangement but absence of somatic hypermutation
immunoglobulin genes of human B cell precursor leukemia.";
J. Exp. Med. 168:229-245(1988).
                                                                                                                                                                                                                                                                                                                  NUCLEOTIDE SEQUENCE.
PubMed=2538551; DOI=10.1084/jem.169.4.1391;
Nickerson K.G., Berman J., Glickman E., Chess L., Alt F.W.;
"Early human IgH gene assembly in Epstein-Barr virus-transform B cell lines. Preferential utilization of the most JH-proximal segment (DQ52) and two unusual VH-related rearrangements.";
J. Exp. Med. 169:1391-1403(1989).
                                                                                                                                                NUCLEOTIDE SEQUENCE.
PubMed=1904154;
Neale G.A., Kitchingman G.R.;
"mRNA transcripts initiating within the human immunoglobulin chain enhancer region contain a non-translatable exon and are extremely heterogeneous at the 5' end.";
Nucleic Acids Res. 19:2427-2433(1991).
     clone: an early step of
  a distinct B lineage?";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    613;
                                                                                                    complementarity-determining regions (CDR3) in human fetal immunoglobulin heavy chain rearrangements."; Eur. J. Immunol. 22:247-251(1992).
                                                                                            and short
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Pred. No. 2.2e-44
; Mismatches 1
                                                                      M.J., Van
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     clone
non of fetal B cells or a 29:1363-1373(1992).
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                                                                      E., Kenter
     "IGM kappa/lambda EBV human B
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InterPro; IPR003597; Ig_c1.
InterPro; IPR003006; Ig_MHC.
InterPro; IPR003596; Ig_v.
Pfam; PF07654; C1-set; 4.
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Matches 104; Conservative
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                                                        PubMed=1730252;
Raaphorst F.M., Timmers
Schuurman R.K.;
"Restricted utilization
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PROSITE; PS50835; IG LI
PROSITE; PS00290; IG ME
Immunoglobulin domain.
SEQUENCE 613 AA; 672
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Mammalia; Butheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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QVQLVESGGGVVQPGRSLRLSCAASGF1FSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
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                                                                                   TISSUE=Rectum tumor;
The German cDNA Consortium;
Wambutt R., Heubner D., Mewes H.W., Weil B., Amid C., Or
Fobo G., Han M., Wiemann S.;
Submitted (JAN-2005) to the EMBL/GenBank/DDBJ databases.
EMBL; BX640627; CAE45781.1; -; mRNA.
HSSP; P01861; 1ADQ.
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                                                                                                                                                                                                                                PRT; 472 AA.

vo-JUL-2004 (TrEMBLrel. 27, Created)

05-JUL-2004 (TrEMBLrel. 27, Last sequence update)

1 05-JUL-2004 (TrEMBLrel. 27, Last annotation update)

Hypothetical protein DKFZp686P15220.

Name=DKFZp686P15220;

Homo sapiens (Human)

Eukarvotta
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InterPro; IPR003597; Ig_c1.
InterPro; IPR003597; Ig_c1.
InterPro; IPR003596; Ig_w.
Pfam; PP07654; C1-set; 3.
SMART; SM00409; IG; 2.
SMART; SM00409; IG; 2.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 4.
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L93 HUMAN
Q9UL93 HUMAN PRELIMINARY;
Q9UL93;
01-MAY-2000 (TrEMBLrel. 13,
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SEQUENCE 472 AA;
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
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Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
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MEDLINE=93301610; PubMed=8315388; DOI=10.1084/jem.178.1.331;
Hillson J.L., Karr N.S., Oppliger I.R., Mannik M., Sasso E.H.;
"The structural basis of germline-encoded VH3 immunoglobulin binding to staphylococcal protein A.";
J. Exp. Med. 178:331-336(1993).
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Bird J., Galili N., Link M., Stites D., Sklar J.;
"Continuing rearrangement but absence of somatic hypermutation immunoglobulin genes of human B cell precursor leukemia.";
J. Exp. Med. 168:229-245(1988).
EMBL; AF035021; AAD56257.1; -; mRNA.
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MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;
Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                fetus.";
Clin. Immunol. Immunopathol. 87:184-192(1998)
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     01-MAR-2004 (TrEMBLrel. 26, Last anno
Myosin-reactive immunoglobulin heavy
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PIR; PH1644; PH1644.

PIR; PL0120; PL0120.

HSSP; P01772; 2FB4.

SMR; Q9UL93; 1-116.

InterPro; IPR007110; Ig-like.

InterPro; IPR003596; Ig_v.

InterPro; IPR003596; Ig_v.
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12434 MW;
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QEGMY2;
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116 AA;
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Last annotation update)
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Pred. No. 2.2e-43;
3; Mismatches 16;
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SMART; SM004007; IGc1; 4.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS00290; IG_MHC; UNKNOWN SEQUENCE 606 AA; 66185 MW; B6i
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NIH MGC Project;
Submitted (JUN-2004) to the EMI
EMBL; BC073758; AAH73758.1; -;
SMR; Q6GMY2; 20-256.
InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_c1.
InterPro; IPR003596; Ig_MHC.
InterPro; IPR003596; Ig_WHC.
InterPro; IPR003596; Ig_WHC.
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Q652C9;
25-OCT-2004 (TrEMBLrel. 28,
25-OCT-2004 (TrEMBLrel. 28,
25-OCT-2004 (TrEMBLrel. 28,
25-OCT-2004 (TrEMBLrel.).
Single-chain Fv (Fragment).
Name=scFv;
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata
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1 Similarity 76.1%;
102; Conservative
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;
Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
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                                                                                                                                                                                                                      Length 240;
                                                       MEDLINE=97362799; PubMed=9219263; DOI=10.1038/nbt0797-629;
MEDLINE=97362799; PubMed=9219263; DOI=10.1038/nbt0797-629;
Kontermann R.E., Wing M.G., Winter G.;
"Complement recruitment using bispecific diabodies.";
"Complement recruitment using bispecific diabodies.";
Nat. Biotechnol. 15:629-631(1997).
EMBL; Y13056; CAA73499.1; -; mRNA.
InterPro; IPR003599; Ig.
InterPro; IPR003596; Ig-v.
SMART; SM00409; IG; 2.
SMART; SM00406; IGv; 2.
PROSITE; PS50835; IG_LIKE; 2.
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
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O9UL90;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2004 (TrEMBLrel. 13, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  complementarity-determining regions (CDR3) in human fetal immunoglobulin heavy chain rearrangements.";

Eur. J. Immunol. 22:247-251(1992).

EMBL; AF035024; AAD56260.1; -; mRNA.

HSSP; P01772; 2FB4.
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Pred. No. 1.7e-43;
6; Mismatches 10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PubMed=1730252;
Raaphorst F.M., Timmers E., Kenter M.J.,
Schuurman R.K.;
"Restricted utilization of germ-line VH3
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                                                                                                                                                                                                 25569 MW;
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ilarity 80.2%;
Conservative
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5: P01772; 2FB4.
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                                         SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                     LVTVSS 117
                                                                                                                                                                                                 240 AA;
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hes 101; Conser
                  NCBI_TaxID=9606;
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                                                   STRAIN=C1q/7
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NUCLECTIDE SEQUENCE.

TISSUE=Primary B-Cells;

MEDINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

A RIAUSENE=2388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

Straubberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Riausner R.D., Collins F.S., Wagner L., Shemmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Altschenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Roback S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Bosak S.A., McEwan P.J., McKernan R.J., Lu X., Gibbs R.A.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

RA Fahey J., Helton B., Ketteman M., Achenko Y., Bouffard G.G.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

Butterfield Y.S.N., Krzywinski W.M., Marra M.A.,

RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.,

"Concert Han 15,000 full-length human M.M., McAndra M.A., McAndra M.A., Marra M.A.,

"Concert Han 15,000 full-length human M.M., McAndra M., McAndra M.A., McAndra M.A., McAndra M.A., Marra M.A.,

"Concert Han 15,000 full-length human M.M., McAndra M.A., Marra M.A., McAndra M.A., McAndra M.A., McAndra M.A., McAndra M.A., McAndra M.A., Marra M.A., McAndra M.A., McAndra M.A., McAndra M.A., McAndra M.A., McHandra M.A., McHandra M.A., McHandra M.A., McHandr
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                     QVQLVESGGGVVQPGRSLRLSCAASGF1FSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                                                                                                                                                                                                                                                                                                                                                                                                                                      Prórbin Sequence.
PubMed=1555592;
Makiya R., Stigbrand T.;
"Placental alkaline phosphatase has a binding site for the human
                                                                                                                                                                                                                                                                                                                                          13;
                                                                                                                                                                                                                                                                                     113;
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Last annotation update)
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
NON_TER 1
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Q8WU38;
01-MAR-2002 (TrEMBLrel. 20,
01-MAR-2004 (TrEMBLrel. 20,
01-MAR-2004 (TrEMBLrel. 26,
IGHD protein.
Name=IGHD;
                                                                                                                                                                                                                                                                                                                                             Conservative
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TISSUE=Primary B-Cells;
Director MGC Project;
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Homo.
NCBI_TaxID=9606;
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21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
19 heavy chain V-III region CAM.
Homo sapiens (Human).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Buarchontoglires; Primates; Catarrhini; Hominidae.
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MEDLINE=81013859; PubMed=6774332;
Lehman D.W., Putnam F.W.;
"Amino acid sequence of the variable region of a human mu chain:
location of a possible JH segment.";
Proc. Natl. Acad. Sci. U.S.A. 77:3239-3243(1980).
-!- MISCELLANEOUS: This mu chain was isolated from the plasma of patient with macroglobulinemia.
-!- SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
                                                                                                                                                                                                                                                                                                                                                                                                    9
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                                                                                                                                                                                                                                                                                                                                                             Score 510; DB 2;
Pred. No. 1.3e-42,
2; Mismatches 13
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HSSP; P01772; 2FB4.
SMR; P01768; 2-122.
GO; GO:0005576; C:extracellular region.
GO; GO:0003823; F:antigen binding; NAS
GO; GO:0006955; P:immune response; NAS
InterPro; IPR007110; Ig-like.
                                                                                                                 sapiens
                                                                                                                                                                                                                                          SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 4.
PROSITE; PS00290; IG MHC; UNKNOWN_2
Immunoglobulin domain; Repeat.
                                      -; mRNA
mmunoglobulin-G Rc portion.";
hr. J. Biochem. 205:341-345(1992)
MBL; BC021276; AAH21276.1; -; mRN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
                                                  PIR; $21205; $21205.

PIR; $30532; $30532.

HSSP; P18529; 118K.

Ensembl; ENSG0000196122; Homo BilnterPro; IPR007110; Ig-like.

InterPro; IPR003597; Ig_c1.

InterPro; IPR003596; Ig_MHC.

InterPro; IPR003596; Ig_W.

Pfam; PF07654; C1-set; 1.

Pfam; PF00047; ig; 2.

SMART; SM00406; IGV; 1.
                                                                                                                                                                                                                                                                                                                                                                                                 12;
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76.6%;
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SEQUENCE 573
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                           multiple
primers."
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NUCLEOTIDE SEQUENCE.

NUCLEOTIDE SEQUENCE.

MEDLINE=96071149; PubMed=7475288;

Cao J., Vescio R.A., Rettig M.B., Hong C.H., Kim A., Lee J.C.,

Lichtenstein A.K., Berenson J.R.;

"A CD10-positive subset of malignant cells is identified in multipl

I myeloma using PCR with patient-specific immunoglobulin gene primers

L Leukemia 9:1948-1953(1995).

R MSD1; S80860; AAD14339.1; -; mRNA.

R EMBL; S80860; AAD14339.1; -; mRNA.

R HSSP; P01842; 1AQK.

R ENSembl; ENSG00000130076; Homo sapiens.

GO; GO:0005887; C:integral to plasma membrane; NAS.

GO; GO:0016066; P:cellular defense response (sensu Vertebrata); NAS InterPro; IPR003110; Ig-like.

R InterPro; IPR003596; Ig_v.
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InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Direct protein sequencing; Immunoglobulin domain;
Immunoglobulin V region; Pyrrolidone carboxylic acid
Immunoglobulin V region; Ig-like.
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Last sequence update)
Last annotation update)
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                                                                                                                                                              Score 507; DB 1;
Pred. No. 4.5e-43;
; Mismatches 11
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Pred. No. 1.4e-42;
7; Mismatches 2:
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13668 MW;
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76.2%;
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illarity 76.2%;
Conservative
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PROSITE; PS50835; IG_LIKE;
                                                                                                                                                                            Similarity 76.3
96; Conservative
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Q9Y509 HUMAN PRELIMINARY
Q9Y509;
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01-NOV-1999 (TrEMBLrel
01-OCT-2003 (TrEMBLrel
VH3 protein (Fragment)
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Homo sapiens (Human)
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122 AA;
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Best Local Similarity
Matches 96; Conser
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MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Riausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Raha S.S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Nillalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Nillalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Nillalon B., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

A Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;

T "Generation and initial analysis of more than 15,000 full-length human
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                                                                                                                                                                                                                                                                Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
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TISSUE=Primary B-Cells;
NIH MGC Project;
Submitted (DEC-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; BC041037; AAH41037.1; -; mRNA.
HSSP; P01861; 1ADQ.
R InterPro; IPR003599; Ig.
R InterPro; IPR003597; Ig.cl.
InterPro; IPR003596; Ig.MHC.
InterPro; IPR003596; Ig.WHC.
InterPro; IPR003596; Ig.V.
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17BED38D917970D6 CRC64;
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Pred. No. 7.8e-42;
); Mismatches 19;
                                                                                                                                                                  Created)
Last sequence up
Last annotation
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Best Local Similarity 76.0%;
Matches 98; Conservative
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27,
27,
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QEPIBL;
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126
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121 TVTVSS
                                    121 LVTVSS
                                                                                                                                                                                                                            IGHM protein.
Name=IGHM;
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05-JUL-2004
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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
80 VDSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCAREFESTMTTVNADYYYFYMDVWG
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PDB; 2FB4; X-ray; H=2-126.

PDB; 2IG2; X-ray; H=2-126.

RG0; G0:0005576; C:extracellular region; NAS.

RG0; G0:0006955; P:immune response; NAS.

RG0; G0:0006955; P:immune response; NAS.

InterPro; IPR007110; Ig-like.

InterPro; IPR003596; Ig_v.

R SMART; SM00406; IGv; 1.

R PROSITE; PS50835; IG_LIKE; 1.

PROSITE; PS50835; IG_LIKE; 1.

A 3D-structure; Direct protein sequencing; Immunoglobulin domain; Immunoglobulin V region; Pyrrolidone carboxylic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=83289131; PubMed=6884994;
Schmidt W.E., Jung H.-D., Palm W., Hilschmann N.;
"Three-dimensional structure determination of antibodies. Primary structure of crystallized monoclonal immunoglobulin IgG1 KOL, I.", Hoppe-Seyler's Z. Physiol. Chem. 364:713-747(1983).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        X-RAY CRYSTALLOGRAPHY (1.9 ANGSTROMS).

MEDLINE=81072295; PubMed=7441755;

Marquart M., Deisenhofer J., Huber R., Palm W.;

"Crystallographic refinement and atomic models of the intact immunoglobulin molecule Kol and its antigen-binding fragment at and 1.0-A resolution.";

J. Mol. Biol. 141:369-391(1980).

-! SIMILARITY: Contains 1 Ig-like (immunoglobulin-like) domain.
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Pyrrolidone carboxylic
                                                                                                                                                                                   21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Ig heavy chain V-III region KOL.
Homo sapiens (Human)
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KGTTVTVSS
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NUCLEOTIDE SEQUENCE.
TISSUB=Primary B-Cells;
NIH MGC Project;
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NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                              Acad.
                                                                                                                                               NCBI_TaxID=9606;
              TVTVSS
                                                                                                                                                                                                                                                                                                             and mouse cDNA |
Proc. Natl. Ace |
                           LVTVSS
                                                                                                                                                                                                                                                                                                                                                                                        PubMed=2500644;
                                                                                                    IGHM protein.
Name=IGHM;
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HUMAN
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                                                                                                                                                 ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARDRGIAAGGNYYYYGMDVWGQGT
                                                                                                                                                                                                                                                                                         Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Euarchontoglires; Primates; Catarrhini; Hominidae
                                                                                                                    QVQLVESGGGVVQPGRSLRLSCAASGFIFSSYAMHWVRQAPGNGLEWVAFMSYDGSNKKY
                                                                                                                           Young D.C.; "Myosin-reactive autoantibodies in rheumatic carditis and normal
                                                                                                    ö
                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE.
NUCLEOTIDE SEQUENCE.
MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;
Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 122;
                                                                                     Length 126;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
                                                                                                    19;
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                                                                        E4D71B52B16F8776
                                                                                     ore 493; DB 1;
red. No. 1.2e-41;
Mismatches 19
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Pred. No. 7.3e-41;
6; Mismatches 13
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-; mRNA.
                                                                                                                                                                                                                                 122 AA
                                                                                      73.8%; Score 493; 76.2%; Pred. No. 1
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Ensembl; ENSG00000130076; Homo sapiens.

InterPro; IPR007110; Ig-like.

InterPro; IPR003596; Ig_v.
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EMBL; AF035030; AAD56266.1; -HSSP; P01772; 2FB4.
SMR; Q9UL84; 1-122.
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illarity 77.0%;
Conservative (
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Q9UL84;
01-MAY-2000 (TrEMBLrel. 13,
01-MAY-2000 (TrEMBLrel. 13,
01-OCT-2003 (TrEMBLrel. 25,
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PROSITE; PS50835; IG_LIKE;
                                                                                                     Conservative
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122 AA;
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96; Conser
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X Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Altachul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gubbs R.A.,

A Nilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

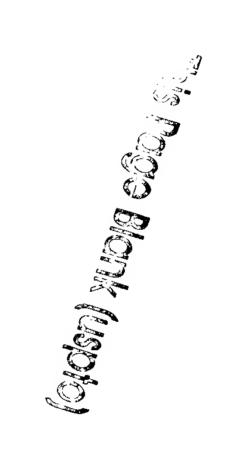
A Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Butterfield Y.S.N., Krzywinski M.I., Marra M.A.;

"Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
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15,000 full-length human
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae
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... Natl. Acad. Sci. U.S.A. 99:16899-16903(2002)
             DB 2;
                                           16;
            Score 484.5; DB 2
Pred. No. 5.2e-40;
; Mismatches 16
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NIH MGC Project;
Submitted (DEC-2001) to the E
EMBL; BC019046; AAH19046.1; -
HSSP; P01861; 1ADQ.
SMR; Q6PJ95; 20-473.
InterPro; IPR003599; Ig.
InterPro; IPR003110; Ig-like.
InterPro; IPR003597; Ig_c1.
InterPro; IPR003596; Ig_wHC.
InterPro; IPR003596; Ig_wHC.
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Pfam; PF07654; C1-8et; 3.
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Name=IGHG1;
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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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MEDLINE=74175307; PubMed=4208843;
Florent G., Lehman D., Putnam F.W.;
Florent G., Lehman D., Putnam F.W.;
"The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains of human IgM immunoglobulins "The switch point in mu heavy chains "The switch point in mu heavy chains" "The switch point in mu heavy chains "The switch point "The switch "The switch point "The switch "The swit
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HSSP; P01772; 2FB4.
SMR; P01769; 5-122.
GO; GO:0005576; C:extracellular region; NAS.
GO; GO:0006955; P:immune response; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig.v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Direct protein sequencing; Immunoglobulin domain;
Immunoglobulin V region; Pyrrolidone carboxylic acid.
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PS50835; IG_LIKE; 4.
PS00290; IG_MHC; UNKNOWN 2.
544 AA; 60102 MW; 1895814B2297C668
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21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Ig heavy chain V-III region GA.
Homo sapiens (Human).
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      Sequence
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US-09-472-087-80
US-08-862-124-17
US-09-269-332-89
US-09-260-198A-2
US-09-240-274-139
US-09-240-274-139
US-09-848-798-139
US-09-840-459-77
US-09-497-625A-77
US-09-497-625A-79
US-09-315-926A-80
US-09-240-274-145
US-09-240-274-16
US-09-848-798-16
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Pred. No. 8.1e
1; Mismatches
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Sequence 70, Application US/09472087

Patent No. 6682736

GENERAL INFORMATION:

APPLICANT: HANSON, DOUGLAS C.

APPLICANT: MUELLER, EILLEN E.

APPLICANT: HANKE, JEFFREY H.

APPLICANT: HANKE, JEFFREY H.

APPLICANT: GILMAN, STEVEN C.

APPLICANT: GORVALAN, JOSE R.

TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBOL

FILE REFERENCE: ABX-PF1

CURRENT FILING DATE: 1999-12-23

PRIOR FILING DATE: 1998-12-23

NUMBER OF SEQ ID NOS: 147

SOFTWARE: PATENTIN Ver. 2.1

SEQ ID NO 70

LENGTH: 451

TYPE: DEM
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Sequence 26, Application US/09240274
Patent No. 6255455
GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QVQLVESGGGVVQPGRSLRLS(
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         sapien
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US-09-848-798-26

Sequence 26, Application US/09848798

Patent No. 6858719

GENERAL INFORMATION:

APPLICANT: Slegel, Donald L.

TITLE OF INVENTION: RA(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

FILE REFERENCE: 09596-42U2

CURRENT APPLICATION NUMBER: US/09/848,798

CURRENT FILING DATE: 2001-05-04

PRIOR APPLICATION NUMBER: EARLIER PILING DATE: 1999-01-29

PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11

NUMBER OF SEQ ID NOS: 224

SEQ ID NO 26

LENGTH: 126
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SORTING METHOD FOR PRODUCTION
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Pred. No. 1e-48;
3; Mismatches
                                                                                                                                                                                                                                                                                                        Score 565; DB 2
Pred. No. 1e-48;
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                                  US/09/240,274
           FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/09/240,274
CURRENT FILING DATE: 1999-01-29
EARLIER APPLICATION NUMBER: 60/081,380
EARLIER FILING DATE: 1998-04-10
EARLIER APPLICATION NUMBER: 60/028,550
EARLIER FILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
SOFTWARE: Patentin Ver: 2.0
                                                                                                                                                                                                                                                     OTHER INFORMATION: anti-Rh(D) chain D31-09-240-274-26
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ilarity 84.1%;
Conservative
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US-09-848-798-26
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106; Conserv
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INVENTION:
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LENGTH: 126
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                                                                                   US-09-240-274-153

i Sequence 153, Application US/09240274

j Patent No. 6255455

j GENERAL INFORMATION:

j TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL

j TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

j TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

j TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

j CURRENT APPLICATION NUMBER: US/09/240,274

j CURRENT FILING DATE: 1999-01-29

j EARLIER PILING DATE: 1998-04-10

j EARLIER PILING DATE: 1996-10-11

j NUMBER OF SEQ ID NOS: 224

j SOFTWARE: Patentin Ver. 2.0
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Sequence 153, Application US/09848798
Patent No. 6858719
GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/09/848,798
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PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER:
PRIOR FILING DATE: EARLIER FILING DATE: 1999-01-29
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER:
PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
SOFTWARE: Patentin Ver. 2.0
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No. 4.1e-48;
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Pred. No. 4.1e-48
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Mismatches
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nilarity 83.3%;
Conservative
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 105; Conser
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Best Local Similarity
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US-09-240-274-153
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OTHER INFORMATION:
09-848-798-153
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ENGTH: 126
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LENGTH: 126
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Sequence 11, Application US/09079029
Patent No. 6342369
GENERAL INFORMATION:
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Ashkenazi, Avi J.
Chuntharapai, Anan
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TELECOMMUNICATION INFORMATION
TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881
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South San Francisco
California
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Best Local Similarity 85.7%;
Matches 108; Conservative
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NAME: Marschang, Diane L
REGISTRATION NUMBER: 35,
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11 Similarity 83.3%;
105; Conservative
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Amino Acid
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APPLICANT: Kim, Kyung J.
TITLE OF INVENTION: Apo-2
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
 2.0
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APPLICATION NUMBER: US
FILING DATE:
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SOFTWARE: Patentin Ver.
SEQ ID NO 25
LENGTH: 126
TYPE: PRT
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                                                                                   OTHER INFORMATION:
-09-848-798-25
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STREET: 1 1
CITY: South
                                                           ORGANISM: Homo
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US-09-079-029-11
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APPLICANT:
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Patent No. 6255455
GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/09/240,274
CURRENT FILING DATE: 1999-01-29
EARLIER APPLICATION NUMBER: 60/081,380
EARLIER PILING DATE: 1998-04-10
EARLIER PILING DATE: 1998-04-10
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TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVITIE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/09/848,798
CURRENT FILING DATE: 2001-05-04
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/240,274
PRIOR FILING DATE: EARLIER FILING DATE: 1999-01-29
PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11
PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11
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Pred. No. 6.4e-48;
Mismatches 13
     Mismatches
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NUMBER OF SEQ ID NOS: 224
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 25
LENGTH: 126
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ilarity 83.3%;
Conservative
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Best Local Similarity
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US-09-240-274-25
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                             Length 126;
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                                                  13;
                             Score 557; DB 2;
Pred. No. 6.4e-48;
Mismatches 13
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Pred. No. 3.6e-47
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IBM PC compatible
YSTEM: PC-DOS/MS-DOS
chain D30
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ORGANISM: Homo sapiens
    ; ORGANISM: Homo sapi
; FEATURE:
; OTHER INFORMATION:
US-09-848-798-140
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US-10-330-613A-29
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-09-424-840B-6
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LENGTH: 120
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; Sequence 140, Application US/09848798
; Patent No. 6858719
; GENERAL INFORMATION:
; APPLICANT: Siegel, Donald L.
; TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
; TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
; FILE REFERENCE: 09596-42U2
; CURRENT APPLICATION NUMBER: US/09/848,798
; CURRENT FILING DATE: 2001-05-04
; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/028,550
; PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11
; NUMBER OF SEQ ID NOS: 224
; SOFTWARE: Patentin Ver. 2.0
                                                     40 QVQLVQSGGGVVQPGRSLRLSCAASGFIFSSYGMHWVRQAPGKGLEWVAGIFYDGGNKYY
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Sequence 140, Application US/09240274

Patent No. 6255455

GENERAL INFORMATION:

APPLICANT: Siegel, Donald L.

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

FILE REFERENCE: 09596-42U2

CURRENT APPLICATION NUMBER: US/09/240,274

CURRENT FILING DATE: 1999-01-29

EARLIER APPLICATION NUMBER: 60/081,380

EARLIER FILING DATE: 1996-04-10

EARLIER FILING DATE: 1996-10-11

NUMBER OF SEQ ID NOS: 224

SOFTWARE: PatentIN Ver. 2.0

SEQ ID NO 140

LENGTH: 125

TYPE: PRT

ORGANISM: Homo sapiens

FEATURE:

FEATURE:

CONTINE:

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Pred. No. 4.5e-47;
9; Mismatches 11;
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US-09-240-274-140
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nilarity 83.3%;
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Best Local Similarity
Matches 105; Conser
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US-09-240-274-140
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; Patent No. 6790938
; GENERAL INFORMATION:
; APPLICANT: Berchtold, Peter
; APPLICANT: Escher, Robert F. A.
; TITLE OF INVENTION: ANTI-GPIIB/IIIA RECOMBINANT ANTIBODIES
; FILE REFERENCE: 100564-09049
; CURRENT APPLICATION NUMBER: US/09/424,840B
; CURRENT FILING DATE: 1999-12-03
; PRIOR APPLICATION NUMBER: DE 19820663.1
; PRIOR FILING DATE: DE 1975527.7
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                                                Score 548.5; DB 2;
Pred. No. 4.5e-47;
9; Mismatches 11;
SH1
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Pred. No. 7.5e-4;
Mismatches
clone
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APPLICANT: Gudas, Jean
TITLE OF INVENTION:
FILE REFERENCE: ABGENIX.022A; CURRENT APPLICATION NUMBER: US/10/330,613A; CURRENT FILING DATE: 2002-12-26; PRIOR APPLICATION NUMBER: 60/346299; PRIOR FILING DATE: 2001-12-18; NUMBER OF SEQ ID NOS: 90; SOFTWARE: FastSEQ for Windows Version 4.0
antibody
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anti-Rh(D)
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Best Local Similarity 87.1%;
Matches 108; Conservative
                                                 ch 82.1%;
1 Similarity 83.3%;
105; Conservative
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; ORGANISM: Homo US-09-424-840B-16
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Sequence 13, Application US/09240274

Patent No. 6255455

GENERAL INFORMATION:

TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

FILE REFERENCE: 09596-42U2

CURRENT APPLICATION NUMBER: US/09/240,274

CURRENT PILING DATE: 1999-01-29

EARLIER APPLICATION NUMBER: 60/028,550

EARLIER PILING DATE: 1996-10-11

NUMBER OF SEQ ID NOS: 224

SOFTWARE: PATENTIN Ver. 2.0

SEQ ID NO 13

LENGTH: 126
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Pred. No. 2.2e-46;
; Mismatches 11
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Pred. No. 4.9e-46
9; Mismatches 1:
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PRIOR FILING DATE: 1997-12-12
PRIOR APPLICATION NUMBER: DE 19
PRIOR FILING DATE: 1997-06-06
NUMBER OF SEQ ID NOS: 128
SOFTWARE: Patentin version 3.1
SEQ ID NO 6
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84.1%;
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les 105; Conservative
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                                                                                       TYPE: PRT ORGANISM: Homo sapiens
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US-09-848-798-13
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GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACT
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
TITLE OF INVENTION SORTING METHOD FOR PRODUCTION THEREOF
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
TITLE OF INVENTION NUMBER: US/09/848,798
CURRENT APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/240,27
PRIOR FILING DATE: EARLIER FILING DATE: 1999-01-29
PRIOR FILING DATE: EARLIER FILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
SOFTWARE: PATENTIN Ver. 2.0
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GENERAL INFORMATION:
APPLICANT: Berchtold, Peter
APPLICANT: Berchtold, Peter
APPLICANT: Berchtold, Peter
TITLE OF INVENTION: ANTI-GPIIB/IIIA RECOMBINANT ANTIBODIES
FILE REFERENCE: 100564-09049
CURRENT APPLICATION NUMBER: US/09/424,840B
CURRENT FILING DATE: 1999-12-03
PRIOR APPLICATION NUMBER: DE 19820663.1
PRIOR FILING DATE: 1998-05-08
PRIOR APPLICATION NUMBER: DE 19755227.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 126;
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Mismatches 12,
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Pred. No. 4.9e-46
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Patent No. 6790938
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PRIOR APPLICATION NUMBER: DE 197
PRIOR FILING DATE: 1997-06-06
NUMBER OF SEQ ID NOS: 128
SOFTWARE: Patentin version 3.1
SEQ ID NO 16
LENGTH: 124
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llarity 82.7%;
Conservative
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Local Similarity 82.7%;
Les 105; Conservative
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ORGANISM: Homo sapiens
FEATURE:
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Qy 120 TTVTVS 126

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Db 118 TTVTVS 124

Search completed: December 28, 2005, 14:05:14

Job time: 36 secs
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US-10-371-942-82

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1731, Ap
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US-10-153-382-17

US-10-153-368-17

US-10-776-649-70

US-11-085-368-17

US-10-77-155-34

US-10-269-711-11

US-10-269-711-23

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US-10-292-088-117
Sequence 117, Application US/10292088
Publication No. US20030211100A1
GENERAL INFORMATION:
APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD F.
APPLICANT: GLADUE, RONALD F.
TITLE OF INVENTION: ANTIBODIES TO CD40
TITLE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/10/292,088
CURRENT APPLICATION NUMBER: 60/348,980
FRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PATENTIN VOR: 2.1
SEQ ID NO 117
LENGTH: 123
TYPE: PRT
ORGANISM: HOMO err
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RESULT 2
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; Publication No. US20030148463A1
; GENERAL INFORMATION:
; APPLICANT: KUFER, PETER
; APPLICANT: RAUM, TOBIAS
; TITLE OF INVENTION: RECEPTORS AND USE

RESULT US-10-3

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APPLICANT: Bowdish, Katherine S.
APPLICANT: Wild, Martha A.
APPLICANT: Wild, Martha A.
APPLICANT: Wild, Martha A.
APPLICANT: Maruyama, Toshiaki
APPLICANT: Maruyama, Toshiaki
APPLICANT: No. US20040009178Alan, Mary Jean
TITLE OF INVENTION: IMMUNOTHERAPEUTICS FOR BIODEFENSE
FILE REFERENCE: 84 (1087-73)
CURRENT APPLICATION NUMBER: US/10/364,743
CURRENT FILING DATE: 2003-02-11
PRIOR APPLICATION NUMBER: US 60/428,807
PRIOR FILING DATE: 2002-11-25
NUMBER OF SEQ ID NOS: 118
SOFTWARE: Patentin version 3.2
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Pred. No. 3.5e-48;
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Pred. No. 4.8e-48;
1; Mismatches 11
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LOCATION: (26)...(26)
OTHER INFORMATION: xaa=unknown amino acid
CURRENT APPLICATION NUMBER: US/10/325,694
CURRENT FILING DATE: 2002-12-19
PRIOR APPLICATION NUMBER: US/09/403,107
PRIOR FILING DATE: 1999-10-14
NUMBER OF SEQ ID NOS: 152
SOFTWARE: Patentin Ver. 2.1
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; ORGANISM: HUMAN
US-10-325-694-144
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ORGANISM: human
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                                                                                  SOFTWARE: Pat
SEQ ID NO 144
LENGTH: 138
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APPLICANT: Frederickson, Shana
APPLICANT: Wild, Martha A.
APPLICANT: Wild, Martha A.
APPLICANT: Maruyama, Toshiaki
APPLICANT: Maruyama, Toshiaki
APPLICANT: Nolan, Mary Jean
TITLE OF INVENTION: IMMUNOTHERAPEUTICS FOR BIODEFENSE
FILE REFERENCE: 98 CIP (1087-73 CIP)
CURRENT APPLICATION NUMBER: US/10/452,593
CURRENT FILING DATE: 2003-06-02
PRIOR APPLICATION NUMBER: US 60/356,086
PRIOR FILING DATE: 2002-02-11
PRIOR FILING DATE: 2002-02-11
PRIOR FILING DATE: 2002-04-29
PRIOR FILING DATE: 2002-04-29
PRIOR FILING DATE: 2002-11-25
                                                                                                              APPLICANT: Frederickson, Shana
APPLICANT: Wild, Martha A.
APPLICANT: Wild, Martha A.
APPLICANT: Waruyama, Toshiaki
APPLICANT: Maruyama, Toshiaki
APPLICANT: No. US20040009178Alan, Mary Jean
TITLE OF INVENTION: IMMUNOTHERAPEUTICS FOR BIODEFENSE
FILE REFERENCE: 84 (1087-73)
CURRENT APPLICATION NUMBER: US/10/364,743
CURRENT FILING DATE: 2003-02-11
PRIOR APPLICATION NUMBER: US 60/428,807
PRIOR FILING DATE: 2002-11-25
NUMBER OF SEQ ID NOS: 118
SOFTWARE: Patentin version 3.2
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Pred. No. 4.8e-48;
4; Mismatches 11
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LOCATION: (26)..(26)
OTHER INFORMATION: xaa=unknown amino acid
0-364-743-98
Sequence 98, Application US/10364743
Sequence 98, Application US/10364743
Publication No. US20040009178A1
GENERAL INFORMATION:
APPLICANT: Bowdish, Katherine S.
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Publication No. US20040258699A1
GENERAL INFORMATION:
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Shana
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PRIOR APPLICATION NUMBER: US 6
PRIOR FILING DATE: 2002-11-25
NUMBER OF SEQ ID NOS: 118
SOFTWARE: PatentIn version 3.2
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Best Local Similarity 86.7%;
Matches 111; Conservative
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ORGANISM: human
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US-10-452-593-14
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SOFTWARE: Pat
SEQ ID NO 150
LENGTH: 138
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US-10-325-694-1
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APPLICANT: Nolan, Mary Jean
TITLE OF INVENTION: IMMUNOTHERAPEUTICS FOR BIODEFENSE
FILE REFERENCE: 98 CIP (1087-73 CIP)
CURRENT APPLICATION NUMBER: US/10/452,593
CURRENT FILING DATE: 2003-06-02
PRIOR APPLICATION NUMBER: US 60/356,086
PRIOR FILING DATE: 2002-02-11
PRIOR APPLICATION NUMBER: US 60/356,086
PRIOR FILING DATE: 2002-02-11
PRIOR FILING DATE: 2002-04-29
PRIOR FILING DATE: 2002-04-29
PRIOR FILING DATE: 2002-04-29
PRIOR FILING DATE: 2002-04-29
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Pred. No. 4.8e-48
                                                                                                         Score 583; DB 5;
Pred. No. 4.8e-48
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; LOCATION: (26)...(26)
; OTHER INFORMATION: xaa=unknown amino acid
US-10-452-593-98
                                                                   amino acid
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Publication No. US20040258699A1
GENERAL INFORMATION:
APPLICANT: Bowdish, Katherine S.
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                           ; FEATURE:
; NAME/KEY: MISC_FEATURE
; LOCATION: (26)..(26)
; OTHER INFORMATION: xaa=unknown
US-10-452-593-14
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ckson, Shana
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illarity 86.7%;
Conservative
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nilarity 86.7%;
Conservative
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NUMBER OF SEQ ID NOS: 118
SOFTWARE: PatentIn version 3.2
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TYPE: PRT
ORGANISM: human
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Best Local S
Matches 111
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APPLICANT:
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Best Local (
Matches 11:
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APPLICANT: RAUM, TOBIAS
TITLE OF INVENTION: NOVEL METHOD FOR THE PRODUCTION OF ANTI-HUMAN ANTIGEN
TITLE OF INVENTION: RECEPTORS AND USES THEREOF
FILE REFERENCE: 38164000
CURRENT APPLICATION NUMBER: US/10/325,694
CURRENT FILING DATE: 2002-12-19
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                                                                      US-10-371-942-82

Sequence 82, Application US/10371942

Publication No. US20030223994A1

GENERAL INFORMATION:

APPLICANT: Hoogenboom, Henricus Renerus Jacobus Mattheus

TITLE OF INVENTION: MHC-PEPTIDE COMPLEX BINDING LIGANDS

FILE REFERENCE: 10280-034001

CURRENT APPLICATION NUMBER: US/10/371,942

CURRENT FILING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: US 60/358,994

PRIOR FILING DATE: 2002-02-20
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Pred. No. 6.8e-48;
5; Mismatches 10;
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Pred. No. 5.5e-48
; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US/09/403,107
PRIOR FILING DATE: 1999-10-14
NUMBER OF SEQ ID NOS: 152
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 121
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 82
LENGTH: 124
TYPE: PRT
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5. US20030148463A1
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11 Similarity 88.9%;
112; Conservative
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il Similarity 87.4%;
111; Conservative
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US-10-371-942-82
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GTTVTVSS
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Publication No. US20
GENERAL INFORMATION:
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US-10-041-860-209
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LENGTH: 126
Query Match
Best Local S
Matches 111
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CANT: Yang, Xiao-Dong
CANT: Yang, Xiao-Dong
CANT: Chen, Francine
CANT: Gazit, Gadi
CANT: Weber, Richard
CANT: Bezabeh, Binyam
OF INVENTION: ANTIBODIES DIRECTED TO PDGFD AND USES
OF INVENTION: THEREOF
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Pred. No. 2.5e-47;
5; Mismatches 10;
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APPLICANT: PRANG, NADJA
APPLICANT: QUADT, CORNELIA
TITLE OF INVENTION: Anti-Epcam Immunoglobulins
FILE REFERENCE: DEBE:031US
CURRENT APPLICATION NUMBER: US/10/778,915
CURRENT FILING DATE: 2004-02-13
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CURRENT APPLICATION NUMBER: US/10/041,860
CURRENT FILING DATE: 2002-01-07
CURRENT FILING DATE: 377
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SOFTWARE: FastSEQ for Windows Version
EQ ID NO 17
LENGTH: 126
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Publication No. US20050180979A1
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ORGANISM: Artificial Sequence
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Jia, Xiao-Chi
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SOFTWARE: PatentIn version
SEQ ID NO 1
LENGTH: 457
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 Length 126;
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APPLICANT: Feng, Xiao-Dong
APPLICANT: Yang, Xiao-Dong
APPLICANT: Chen, Francine
APPLICANT: Gazit, Gadi
APPLICANT: Weber, Richard
APPLICANT: Weber, Richard
APPLICANT: Hereof
TITLE OF INVENTION: THEREOF
FILE REFERENCE: ABGENIX.051A
CURRENT APPLICATION NUMBER: US/10/041,860
CURRENT FILING DATE: 2002-01-07
NUMBER OF SEQ ID NOS: 377
SOFTWARE: FastSEQ for Windows Version 4.0
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 Score 581; DB 4;
Pred. No. 6.9e-48
; Mismatches 1:
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Pred. No. 6.9e-48
3; Mismatches 13
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Publication No. US20030157109A1
GENERAL INFORMATION:
APPLICANT: Corvalan, Jose R.F.
APPLICANT: Jia, Xiao-Chi
APPLICANT: Feng, Xiao
APPLICANT: Yang, Xiao-Dong
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APPLICANT: Jia, Xiao-Chi
APPLICANT: Feng, Xiao
APPLICANT: Yang, Xiao-Dong
APPLICANT: Chen, Francine
APPLICANT: Gazit, Gadi
APPLICANT: Weber, Richard
87.0%;
ilarity 88.1%;
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illarity 88.1%;
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Jia, Xiao-Chi
Feng, Xiao
Yang, Xiao-Dong
Chen, Francine
Gazit, Gadi
Weber, Richard
Bezabeh, Binyam
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US-10-041-860-209
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US-10-292-088-116
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APPLICANT: LaRochelle, William
APPLICANT: Lichenstein, Henri
TITLE OF INVENTION: METHOD FOR THE TREATMENT OF NEPHRITIS
TITLE OF INVENTION: USING ANTI-PDGF-DD ANTIBODIES
FILE REFERENCE: ABGENIX.052A
CURRENT APPLICATION NUMBER: US/10/665,383
CURRENT FILING DATE: 2003-09-16
PRIOR APPLICATION NUMBER: 60/411,137
PRIOR FILING DATE: 2002-09-16
NUMBER OF SEQ ID NOS: 97
SOFTWARE: FastSEQ for Windows Version 4.0
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  INVENTION: ANTIBODIES DIRECTED TO PDGFD AND USES
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6.9e-48;
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Pred. No. 6.9e-48
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Pred. No. 6.9e-
3; Mismatches
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Mismatches
TITLE OF INVENTION: ANTIBODIES DIRECTED TY
TITLE OF INVENTION: THEREOF
FILE REFERENCE: ABGENIX.051A
CURRENT APPLICATION NUMBER: US/10/041,860
CURRENT FILING DATE: 2002-01-07
NUMBER OF SEQ ID NOS: 377
SOFTWARE: FastSEQ for Windows Version 4.0
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Publication No. US20040141969A1
GENERAL INFORMATION:
APPLICANT: Floege, Juergen
APPLICANT: Gazit, Gadi
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illarity 88.1%;
Conservative
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LENGTH: 126
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Pred. No. 7.5e-48;
; Mismatches 9;
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No. 1.1e-47
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                                                                                                                                TITLE OF INVENTION: ANTIBODIES TO CD40
FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/10/292,088
CURRENT FILING DATE: 2003-03-14
PRIOR APPLICATION NUMBER: 60/348,980
PRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
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Sequence 116, Application US/10292088 Publication No. US20030211100A1 GENERAL INFORMATION:
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5. US20030211100A1
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PRIOR APPLICATION NUMBER: 60/348,980
PRIOR FILING DATE: 2001-11-09
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APPLICANT: GLADUE, RONALD P.
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
TITLE OF INVENTION: ANTIBODIES TO CI
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88.9%;
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Best Local Similarity 89.7%;
Matches 113; Conservative
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                                              APPLICANT: BEDIAN, VAHE
APPLICANT: GLADUE, RONALD P
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
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0-292-088-116
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Best Local Similarity
Matches 112; Conser
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LENGTH: 123
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Db 61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGHQLL--GYYYYYGMDVWGQGT 118

Qy 121 TVTVSS 126

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Minimum Match 0% Maximum Match 100% Listing first 45 summaries seq length: 0 seq length: 200000000 Post-processing: Minimum DB Maximum DB

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# SUMMARIES

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### ALIGNMENTS

Tumour necrosis factor alpha; TNF; antibody; light chain; CDR; complementarity determining region; antirheumatic; antiarthritic; antiulcer; antiasthmatic; antiallergic; antiinflammatory; antisickling; antidiabetic; antiarteriosclerotic; antiatherosclerotic; vasotropic; antianginal; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy. antibody light chain variable region. Z standard; protein; 108 Anti-tumour necrosis factor entry) (first 0-JUN-2002 AAM51165; AAM51165 RESULT 1 

Location/Qualifiers 1. .23 Homo sapiens. Key Region

CDR2 CDR3 24. 34 /label= CDR1 35. 49 /label= FR2 50. .56 /label= CDR2 57. .88 /label= FR3 89. .98 /label= CDR3 99. .108 CDR1 ı. .23 /label= FR1 Region Region Region Region Region Region

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2001WO-US024785 07-AUG-2001; 14-FEB-2002

2000US-0223360P. 2000US-0236826P. 2001US-00920137. 07-AUG-2000; 29-SEP-2000; 01-AUG-2001;

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                                                   for
                                                                                                                                                                                                                                                                                                                                      viral, and fungal infections, pneumonia, leprosy and malaria, a malignadisease such as leukaemia, chronic myelocytic leukaemia, Burkitt's lymphoma and multiple myeloma, or a neurological disease such as multipsclerosis, Parkinson's disease, spinal ataxia, Alzheimer's disease and Creutzfeldt-Jakob disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                   useful
                                                             osis,
                                                 Novel isolated mammalian anti-tumor necrosis factor antibody, treating sickle cell anemia, diabetes, atherosclerosis, restenangina pectoris, myocardial infarction, leprosy.
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Pred. No. 1.1e-37;
; Mismatches 1
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          Heavner G,
                                                                                           Claim 9; Page 129-130; 131pp; English.
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                              2002-217194/27
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les 107; Conser
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        Giles-Komar J,
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The present sequence is that of a human Vg/38K-type light chain variable region encoded by a human germline gene in a transgenic mouse used in human monoclonal antibody (mAb) construction. A GenTWV fusion was performed using spleen cells from a hybrid mouse containing human performed using spleen cells from a hybrid mouse containing human performed using spleen cells from a hybrid mouse containing human contains and the splent splent was a lambusised with recombinant human immobilised human TNF alpha with apparently high recombinant human immobilised human TNF alpha with apparently high contained that bound immobilised human TNF alpha with apparently high portion of the light chain variable region of 2 of the mAbb, TWN14 and TWN15, were identical (see AAM51174) to the present sequence, while the light chain variable region of 2 others, TWN148(8) and TWN196 (see AAM51175), differed by a single amino acid residue. The invention remunoglobulist, and cleavage products and variants, as well as anti-TNF mutubolists, immunoglobulist, and cleavage products and variants, as well as anti-TNF complementaring region (CDR) grafted anti-TNF antibodiss, immunoglobulis, and cleavage products and variants, as well as anti-TNF artisegenic plants, and methods of making and using them. The anti-TNF artisegenic plants and methods of making and using them. The anti-TNF artisegenic plants and methods of making and using them. The anti-TNF artisegenic plants and methods of making and using the TNF induced cell adhesion especially the heavy chain and/or light chain variable regions given in molecules, inhibit TNF humanical artibution in a cell, tissue, organ or animal (claimed) such as atterioracies may inhibit TNF induced cell adhesion molecules, inhibit TNF barbanding tor receptor, or provide Arthritic and improvement in a mouse model. They are useful for diagnosing or treating a Theumatoid arthritis, gastric ulcer, asthma, allergic rhining as theumatoid arthritis, gastric ulcer, asthma, allergic chinical viral, and fungal infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel isolated mammalian anti-tumor necrosis factor antibody, useful treating sickle cell anemia, diabetes, atherosclerosis, restenosis, angina pectoris, myocardial infarction, leprosy.
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Location/Qualifiers
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2000US-0236826P
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N-PSDB; ABL53513.
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01-AUG-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; antibody; dual integrin; HC CDR; variable region; LC CDR; medical device; immune related disease; rheumatoid arthritis; gastric ulcer; asthma; allergic rhinitis; Crohn's pathology; sickle cell anaemia; diabetes; cardiovascular disease; arteriosclerosis; atherosclerosis; restenosis; angina pectoris; myocardial infarction; infectious disease; pneumonia; leprosy; malaria; malignant disease; leukaemia; chronic myelocytic leukaemia; multiple myeloma; neurological disease; multiple sclerosis; Parkinson's disease; Alzheimer's disease; Creutzfeldt-Jakob disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel isolated mammalian anti-dual integrin antibody, useful for diagnosing or treating dual integrin related condition such as arthritis, gastric ulcer, asthma, atherosclerosis, restenosis.
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                                                                                                        Length 108;
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and Creutzfeldt-Jakob disease
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                                                                                                       Score 559; DB 5;
Pred. No. 1.1e-37
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Alzheimer's disease
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                                                                                                                                Similarity
                                                    Sequence 108 AA;
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                                                                                                                            Local Sim
hes 107;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAU76334;
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                                                                                                                                                                                                               -
                                                                                                        Query Match
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                                                                                                                                   Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU76334
                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT
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This invention relates to novel mammalian CDR mimetibodies, specific portions or variants thereof. Specifically, it refers to an antibody fragment where a protein has been inserted into, or replaces a portion of, one or more CDR regions, such that each CDR mimetibody comprises at least one portion of a heavy chain or light chain variable region, which itself comprises at least one human framework region and at least one ligand binding region (LBR). The present invention describes human mimetibodies, including modified immunoglobulins and cleavage products
                                                                                                                                                                                                                                                                                                                                             9
sickle cell anaemia, diabetes, cardiovascular disease such as arteriosclerosis, atherosclerosis, restenosis, angina pectoris, myocardial infarction, infectious disease in a cell such as bacterial, viral, and fungal infections, pneumonia, leprosy, malaria; malignant disease such as leukaemia, chronic myelocytic leukaemia, Burkitt's lymphoma, multiple myeloma; neurological disease such as multiple sclerosis, Parkinson's disease, spinal ataxia, Alzheimer's disease, Creutzfeldt-Jakob disease and many other diseases given in the specification. The present sequence is an anti-dual integrin human variable region containing at least one of the six CDRs listed above (AAU76327-AAU76332)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mimetic; CDR mimetibody; gene therapy; transgenic; immune; cardiovascular; infectious; malignant; neurologic disease; anaemia; immunomodulator; cardiant; antimicrobial; cytostatic; neuroprotective; erythropoietin.
                                                                                                                                                                                                                                                                                                                                             1 BIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                                                                                                                                                                1 EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New CDR mimetibody comprising a portion of a heavy or light chain variable region comprising human framework or ligand binding region, useful for preparing a composition for treating e.g., immune, cardiovascular or neurologic disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                    RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
                                                                                                                                                                                                                                                                   Length 108;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                           1;
                                                                                                                                                                                                                                                                     Score 559; DB 5;
Pred. No. 1.1e-37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ghrayeb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SegiD
                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; SEQ ID NO 989; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chain mimetibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scallon BJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADJ73533 standard; protein; 108
                                                                                                                                                                                                                                                                                                         ö
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                                                                                                                                                                                                                                                                98.4%;
99.1%;
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                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (CENZ ) CENTOCOR INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Erythropoietin light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-804237/75.
                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                   Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2003084477-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-MAY-2004
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                                                                                                                                                                                                                                                                                                       107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADJ73533;
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                                                                                                                                                                                                                                                                                                         Matches
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that can be useful in gene therapy and the generation of transgenic plants and animals. Furthermore, the CDR mimetibody is useful for preparing compositions for modulating, treating or reducing the symptoms of immune, cardiovascular, infectious, malignant and/or neurologic diseases, as well as anaemia. Accordingly, they exhibit immunomodulator, cardiant, antimicrobial, cytostatic and neuroprotective activities. This polypeptide sequence is an erythropoietin light chain mimetibody of the
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                                                                                                                                                                                                                                                                                           60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mimetic; CDR mimetibody; gene therapy; transgenic; immune;
cardiovascular; infectious; malignant; neurologic disease; anaemia;
immunomodulator; cardiant; antimicrobial; cytostatic; neuroprotective;
                                                                                                                                                                                                                                                                                           RATGIPA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chain
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                                                                                                                                                                                           Length 108;
                                                                                                                                                                                                                                                                                                                                               RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK
                                                                                                                                                                                                                                                                                                                              RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK
                                                                                                                                                                                                                                                             EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLI
                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chain mimetibody SeqID 990.
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                                                                                                                                                                                         ; DB 7;
1.1e-37;
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                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2; SEQ ID NO 990; 97pp; English.
                                                                                                                                                                                           Score 559;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scallon
                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard; protein; 108
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                                                                                                                                                                                           98.4%;
99.1%;
                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (CENZ ) CENTOCOR INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Erythropoietin light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-804237/75
                                                                                                                                                                                                           Similarity
                                                                                                                                                         Sequence 108 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2003084477-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         erythropoietin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-MAY-2004
                                                                                                                                                                                                                         107;
                                                                                                                            invention.
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Best Local
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ADJ73534
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The invention describes an isolated binding member comprising at least one binding domain capable of specifically binding Streptococcus pneumoniae surface adhesin A (PBaA) protein, the binding domain having a dissociation constant Kd for PBaA which is less than 1 x10 -6 M. Also described are: an isolated nucleic acid molecule encoding at least a part of the binding member described above; a vector comprising the nucleic acid molecule of (1); a host cell comprising the nucleic acid molecule of (1); a cell line engineered to express the binding member described above; detecting or diagnosing a disease or disorder associated with prouncoccus in an individual; a kit comprising at least one binding ac least one binding member described above. The composition comprising at least one binding member described above. The binding member is useful for producing a pharmaceutical composition for the treatment of Pneumococcus infection. The binding member, methods and
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                engineering;
of immune, cardiovascular, infectious, malignant and/ or neurologic diseases, as well as anaemia. Accordingly, they exhibit immunomodulator, cardiant, antimicrobial, cytostatic and neuroprotective activities. This polypeptide sequence is an erythropoietin light chain mimetibody of the invention.
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                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Streptococcus pneumoniae surface adhesin
g or preventing pneumonia, meningitis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antibacterial; antiinflammatory; immunosuppressive; antibody engin
pharmaceutical; infection; pneumonia; meningitis; sepsis;
pneumococcal surface adhesin A; PsaA; light chain variable region.
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                                                                                                                                                                                                                                                               108
                                                                                                                                       Length 108;
                                                                                                                                                                                                                                                             RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK
                                                                                                                                                                       Indels
                                                                                                                                                           ...
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                                                                                                                                      Score 559; DB 7;
Pred. No. 1.1e-37;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; SEQ ID NO 40; 137pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lundgren JD,
                                                                                                                                                                                                                                                                                                                                                                         ADW07091 standard; protein; 108 AA
                                                                                                                                                                       ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ing member towards Stuseful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2004WO-DK000492
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2003US-0486647P
                                                                                                                                         98.4%;
99.1%;
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                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Anti-PsaA-antibody
                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2005-101476,
N-PSDB; ADW07090
                                                                                                             Sequence 108 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-JUL-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
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suffering
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           pneumonia
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                                                                                                                         RATGIPA
                                                                                                      Gaps
                               chain
 individual
composition are useful for treating or preventing an individua from disorders or diseases associated with S. pneumoniae, e.g. meningitis and/or sepsis. This is the amino acid sequence of a pneumococcal surface adhesin A (PsaA)-antibody 9A7 kappa light
                                                                                                      ö
                                                                                                                                     TNF; antibody; heavy chain; CDR; sgion; antirheumatic; antiarthrit
                                                                                                                          EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASN
                                                                                                                                                                                                                                                                                                         region
                                                                                                                                                                  108
                                                                                 Length 108;
                                                                                                                                                                  RFSGSGSGTDFTLTISSLEPEDFAVYYCOORSNWPPFTFGPGTKVDIK
                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                       chain variable
                                                                                           1.1e-37
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                                                                                 DB
                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                      TNV15 light
                                                                                Score 559;
Pred. No. 1
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2000US-0236826P.
2001US-00920137.
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/label= FR2
                                                                                98.4%;
99.1%;
                                                                                                                                                                                                                                          protein;
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/label= FR1
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label= FR3
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/label= M
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                                                                                                      Conservative
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/label=
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119. .12
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M
                              pneumococcal surface variable region.
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                                                                                           al Similarity
107; Conser
                                                                                                                                                                                                                                                                                                         recombinant
                                                             108 AA;
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29-SEP-2000;
01-AUG-2001;
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                                                                                                                                                                                                                                                                                   10-JUN-2002
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                                                              Sequence
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                                                                                  Query Match
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                                                                                                      Matches
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The present sequence is that of the light chain variable region of antitumour necrosis factor (TNP) human recombinant monoclonal antibodies (mAbb) TWV14 and TWV15. These are human mabs produced from a GenTWV containt of the using spleen cells from a hybrid monose containing human TMP alpha. The human mabs bind immobilised human TNP alpha with high avidity and haran is gai, kappa isotype. The mature portion of the light chain variable region is identical to that of the Vg/38-type light chain variable regions is identical to that of the Vg/38-type light chain variable regions (see AAM51173). The light chain variable regions (collang gradied anticollang provides human, primate, mammalian, rodent, chimeric, the antibodies, immunoglobulins, cleavage products and other specified of the invention provides human, primate, mammalian, rodent, chimeric, TWP antibodies, immunoglobulins, cleavage products and other specified compositions, formulations, devices, transgenic animals, transgenic portions and variants, as well as anti-TWP antibody compositions, formulations, devices, transgenic animals, transgenic plants, and methods of maxing and using them. The anti-TMP antibody compositions, formulations, devices, transgenic animals, transgenic plants, and methods of maxing and using them. The anti-TMP antibody compositions, formulations, devices, transgenic animals, transgenic sequence and in AAM51155, or either all of the CDRs of the C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ô
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21 EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     YDASNRATGIPA
                                                                                              antibody, useful
sis, restenosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gарв
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chain; CDR;
antiarthritic;
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  Ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 129;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indela
                                                                                              Novel isolated mammalian anti-tumor necrosis factor anti
treating sickle cell anemia, diabetes, atherosclerosis,
angina pectoris, myocardial infarction, leprosy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TNV196 light chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tumour necrosis factor alpha; TNF; antibody; heavy complementarity determining region; antirheumatic;
 Scallon B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .. H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 559; DB 5;
Pred. No. 1.3e-37
; Mismatches
    Ö
    Heavner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAM51175 standard; protein; 129 AA
                                                                                                                                                                           Example 3; Fig 5; 131pp; English.
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ilarity 99.1%;
Conservative
    Knight DM,
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                                        2002-217194/27.
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                                                           N-PSDB; ABL53514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 129 AA;
٦,
    Giles-Komar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-JUN-2002
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Best Local (
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Matches
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antiulcer; antiasthmatic; antiallergic; antiinflammatory; antisickling; antidiabetic; antiarteriosclerotic; antiatherosclerotic; vasotropic; antianginal; cardiant; antibacterial; virucide; fungicide; antileprotic; protozoacide; cytostatic; neuroprotective; antiparkinsonian; nootropic; human; diagnosis; therapy; TNV148(B); TNV196; monoclonal antibody; mAb.
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### Homo sapiens.

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Signal_peptide
                            Mature protein
Location/Qualifiers
                                                                                    CDR2
                                                                                                                CDR3
                                                        CDR1
                                                                                                 = FR3
118
                                                                      FR2
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      Peptide
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### WO200212502-A2

### 14-FEB-2002

07-AUG-2001; 2001WO-US024785

07-AUG-2000; 2000US-0223360P 29-SEP-2000; 2000US-0236826P 01-AUG-2001; 2001US-00920137

## (CENZ ) CENTOCOR INC.

Giles-Komar J, Knight DM, Heavner G, Scallon B, Shealy D;

### TOO INTO I THE

WPI; 2002-217194/27.

N-PSDB; ABL53515.

Novel isolated mammalian anti-tumor necrosis factor antibody, useful treating sickle cell anemia, diabetes, atherosclerosis, restenosis, angina pectoris, myocardial infarction, leprosy.

for

# Example 3; Fig 5; 131pp; English.

The present sequence is that of the light chain variable region of antitumour necrosis factor (TNF) human recombinant monoclonal antibodies (mAbs) TNV148(B) and TNV196. These are human mAbs produced from a GenTNV fusion using spleen calls from a hybrid mouse containing human variable and constant region antibody transgenes that was immunised with human TNF alpha. The human mAbs bind immobilised human TNF alpha with high avidity and have a totally human 1gG1, kappa isotype. The mature portion of the light chain yermline sequence (see AAM51173) by a single residue (Tyr-50 creplacing Ser). The light chain variable regions of 2 other human mAbs (see AAM51174) were identical to the germline sequence. The invention provides isolated human, primate, rodent, mammalian, chimeric, humanised and/or complementarity determining region (CDR)-grafted anti-TNF antibody compositions, compositions, devices, transgenic animals, transgenic portions and variants, as well as anti-TNF antibody compositions, compositions, formulations, devices, transgenic animals, transgenic plants, and methods of making and using them. The anti-TNF antibody compositions, comprises at least a portion of an immunoglobulin molecule, especially the heavy chain and/or light chain variable regions given in the present sequence and in AAM51165, or either all of the CDRs of the heavy chain (see AAM51158-60) or all of the CDRs of the heavy chain

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                                                                                                                              a malignant
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inhibit TNF binding to receptor, or provide Arthritic Index improvement in a mouse model. They are useful for diagnosing or treating a TNF related condition in a cell, tissue, organ or animal (claimed) such as rheumatoid arthritis, gastric ulcer, asthma, allergic rhinitis, Crohn's pathology, sickle cell anaemia, diabetes, a cardiovascular disease such as arteriosclerosis, atherosclerosis, restenosis, angina pectoris or myocardial infarction, an infectious disease in a cell such as bacterial, viral, and fungal infections, pneumonia, leprosy and malaria, a malignant disease such as leukaemia, chronic myelocytic leukaemia, Burkitt's lymphoma and multiple myeloma, or a neurological disease such as multiple sclerosis, Parkinson's disease, spinal ataxia, Alzheimer's disease and
                                                                                                                                                                                                                                                                                                                       9
                                                                                                                                                                                                                                                                                                                                                     80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New human monoclonal antibody that specifically binds to growth factor receptor HER2/neu, for treating, preventing or diagnosing diseases characterized by aberrant HER2/neu expression e.g. cancers.
                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention provides the protein and coding sequences for be monoclonal antibodies which bind specifically to the HER2/neu growth factor receptor (also known as erbB2). These are designated 3-F2, 1-I and 2-E8. They can be used in the immunotherapy-based treatment and prognosis of cancers, particularly adenocarcinomas such as salivary gland, stomach, kidney, mammary gland, lung and squamous cell carcing and ovarian cancer. The present sequence is part of an antibody of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer; antibody; immunotherapy;
                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                    108
                                                                                                                                                                                                                                                         Length 129;
                                                                                                                                                                                                                                                                                                                                                                                 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK
                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                        Score 559; DB 5; I
Pred. No. 1.3e-37;
); Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2-E8 light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 110; 113pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    growth factor receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oncogene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB72884 standard; protein; 108
                                                                                                                                                                                                                                                                                       ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human anti-HER2/neu antibody
                                                                                                                                                                                                                                                        98.4%;
ilarity 99.1%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JUL-1999; 99US-0146313P
10-MAR-2000; 2000US-0188539P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000WO-US020272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                sclerosis, Parkinson's di
Creutzfeldt-Jakob disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
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N-PSDB; AAF75590.
                                                                                                                                                                                                                                                                      l Similarity
107; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HER2; neu;
                                                                                                                                                                                                                              129 AA;
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                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB72884;
                                                                                                                                                                                                                                                                                                                                                                                                                     81
                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                       61
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of IP-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   which
                                                                                                                                                                                                                                                                                                                                                                                                                 disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (IP-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antibody is
                                                                                                                                           EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         induced
                                                                                                                     EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .g.,
                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunosuppressive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to an isolated human monoclonal antibody or an antigen-binding portion, which specifically binds to human IP-10 and exhibits at least one property selected from: inhibits binding of IF to CXCR3; inhibits IP-10 induced calcium flux; inhibits IP-10 induced react with mouse IP-10; does not cross-react with human ITAC. The isolated human monoclonal antibody useful for treating an inflammatory or autoimmune disease including multiple sclerosis, rheumatoid arthritis, inflammatory bowel disease (e.g., ulcerative colitis, Crohn's disease), systemic lupus erythematosus, Type I diabetes, inflammatory skin disorders (e.g., graves' psoriasis, lichen planus), autoimmune thyroid disease (e.g., Graves'
                                                                                                                                                                                                                                                                                                                                                                                                                             respiratory disease, neurological disease, degeneration; infection; Neuroprotective, Antiarthritic; Antirheumatic; Antiinflammatory; Gastrointestinal-Gen.; Antiulcer; Dermatological; Immunosuppressive, Antidiabetic; Antipsoriatic; Antithyroid; Antiasthmatic; Respiratory-Gen.; Cerebroprotective; Vasotropic; Nootropic; Antiparkinsonian; Antiangiogenic; Antiarteriosclerotic; Virucide; Antibacterial; light chain variable region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated human monoclonal antibody or an antigen-binding portic
specifically binds to human interferon gamma inducible protein
10), useful for treating viral or bacterial infection, or infla
                                                                                                                                                                                                                                                                                                                                                                                                              IP-10; inflammation; immune disorder; dermatological
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ပ်
                                                                                                                                                                                                               108
                                                                                                                                                                                  108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wang
                                                           Length 108;
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                                                                                                                                                                                  RESGSGSGTDFTLTISSLEPEDFAVYYCOORSNWPPFTFGPGTKVDIK
                                                                                                                                                                                                   RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPYTFGQGTKLEIK
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                                                                                        Indels
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                                                                                                                                                                                                                                                                                                                                                                                   SEQ
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Keirstead HS,
                                                                                        7
                                                                                                                                                                                                                                                                                                                                                                                   Human IP10 antibody light chain variable region,
                                                           4.
                                                         ore 541; DB 4;
ed. No. 3e-36;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 90; 179pp; English.
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Lane TE,
                                                           Score
Pred.
                                                                                                                                                                                                                                                                                          Æ
                                                                                                                                                                                                                                                                                          standard; protein; 108
                                                                                        3;
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                                                        95.2%;
illarity 95.4%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                      entry)
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VS,
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autoimmune diseases
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                                                                                                                                                                                                                                                                                                                                                      (first
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                                                                      Local Similarity
nes 103; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB; AEB01088
                             Sequence 108 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                   antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Deshpande
invention
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                                                                                                                                                                                                                                                                                                                        AEB01062;
                                                                                                                                                                                                                61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo
                                                                                        Matches
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                                                                                       inflammatory
disease, Hashimoto's thyroiditis), Sjogren's syndrome, pulmonary inflammation (e.g., asthma, chronic obstructive pulmonary disease, pulmonary sarcoidosis, lymphocytic alveolitis), transplant rejection, spinal cord injury, brain injury (e.g., stroke), neurodegenerative diseases (e.g., Alzheimer's disease, Parkinson's disease), gingivitis, gene therapy-induced inflammation, diseases of angiogenesis, inflammationether disease (e.g., IgA nephropathy, membranoproliferative glomerulonephritis, rapidly progressive glomerulonephritis), or atherosclerosis. It is also useful for treating a viral or bacterial infection involving unwanted IP-10 activity in a subject, where the virilection is mediated by HIV, HCV, HSV-1 or SARS virus. The present sequence represents the amino acid sequence of a human IP10 monoclonal antibody light chain variable region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New isolated antibody that specifically binds osteoprotegerin ligand, useful for diagnosing or treating bone disorders, such as osteoporosis, bone loss from arthritis, Paget's disease or osteopenia.
                                                                                                                                                                                                                                                                                                                                                  EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                                                                                                                                                                                                                                                                       EIVLTOSPATLSLSPGERATLSCRASOSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human; antibody; osteoprotegerin ligand; OPGL; osteopenic disorder; osteopathic; antiarthritic; cytostatic; gene therapy; bone disorder; osteoporosis; bone loss; arthritis; Paget's disease; osteopenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Huang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16E1 anti-OPGL antibody kappa chain variable region SEQ ID NO:24
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                                                                                                                                                                                                                                                                                   Length 108;
                                                                                                                                                                                                                                                                                                                                                                                                                       Martin
                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                  9
                                                                                                                                                                                                                                                                                   Score 541; DB 9
Pred. No. 3e-36;
                                                                                                                                                                                                                                                                                                    ed. No. 3e-3
Mismatches
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                                                                                                                                                                                                                                                                                                                   3;
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                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 95.4%;
Matches 103; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
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B; ADF11410.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 6; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Medlock
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                                                                                                                                                                                                                                                     108 AA;
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has
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sample with (I) under conditions that allow for binding of the antibody to OPGL, and measuring the level of bound antibody in the sample. (I) he osteopathic, antiarthritic and cytostatic activities, and can be used in gene therapy. The composition and methods are useful in diagnosing or treating bone disorders, such as osteoporosis, bone loss from arthritis, Paget's disease or osteopenia. The antibody (I) may also be used for detecting OPGL in biological samples and in identifying cells or tissues that produce the protein. The present sequence represents a sequence which is used in the exemplification of the present invention.
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Sequence 109 AA;

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                                                    1 EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                         Gaps
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                                                                             RESGSGSGTDFTLTISSLEPEDFAVYYCOORSNWPPFTFGPGTKVDIK 108
        Length 109;
                                                                                    Indels
; DB 7; Le
3.1e-36;
      Score 541; DB Pred. No. 3.1e 3; Mismatches
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       95.2%;
95.4%;
                        Conservative
                Local Similarity
                       103;
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        Query Match
                         Matches
                 Best
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ADF11435 standard; protein; 215 entry) (first 12-FEB-2004 ADF11435

human; antibody; osteoprotegerin ligand; OPGL; osteopenic disorder; osteopathic; antiarthritic; cytostatic; gene therapy; bone disorder; osteoporosis; bone loss; arthritis; Paget's disease; osteopenia. chain SEQ ID NO:48 anti-OPGL antibody light 

Homo

WO2003086289-A2

23-OCT-2003

2003WO-US010749 07-APR-2003; 2002US-0370407P 05-APR-2002;

(AMGE-) AMGEN INC

ligand WPI; 2003-845253/78 N-PSDB; ADF11434.

Huang H;

Martin F,

Elliott RL,

Sullivan JK,

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Medlock

Boyle WJ,

eoporosi New isolated antibody that specifically binds osteoprotegerin useful for diagnosing or treating bone disorders, such as ost bone loss from arthritis, Paget's disease or osteopenia. useful

156pp; English SEQ ID NO 48; Claim 16; The present invention describes an isolated human antibody (I) that specifically binds osteoprotegerin ligand (OPGL). Also described: (1) a pharmaceutical composition comprising a pharmaceutical carrier and a therapeutic amount of (I); (2) methods of treating an osteopenic disorder in a patient, comprising administering to a patient the pharmaceutical composition of (1) or a pharmaceutical amount of (I); and (3) a method for detecting OPGL in a biological sample, comprising contacting the sample with (I) under conditions that allow for binding of the antibody to OPGL, and measuring the level of bound antibody in the sample. (I) has osteopathic, antiarthritic and cytostatic activities, and can be used in gene therapy. The composition and methods are useful in diagnosing or

alling by competing with IL-se MAbs, as well as single chain antibodies, Fab' antibodies are used in methods of treating

The present sequence is that of human anti-interleukin-1 receptor type (IL-1R1) monoclonal antibody (MAb) 26F5 kappa chain variable region. Human MAbs to IL-1R1 were prepared using the HCo7 strain of transgenic mice, which expresses human antibody genes. These mice were immunised with purified recombinant IL-1R1, and splenocytes from immunised mice were fused to a mouse myeloma cell line to generate hybridomas. Hybridomas which secreted a MAb that bound with high avidity to IL-1R1 were selected. The MAbs inhibit IL-1 signalling by competing with IL-1 beta and IL-1alpha binding to IL-1R. These MAbs, as well as single chantibodies single chain Fv antibodies, Fab antibodies, Fab' antibodies of treat:

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English.

179pp;

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Claim

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treating bone disorders, such as osteoporosis, bone loss from arthritis, Paget's disease or osteopenia. The antibody (I) may also be used for detecting OPGL in biological samples and in identifying cells or tissues that produce the protein. The present sequence represents a sequence which is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                             region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      binds interleukin-1 receptor mediated diseases such as inflammatory conditions.
                                                                                                                        EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                Human; monoclonal antibody; antibody; interleukin-1; receptor; antiasthmatic; antiinflammatory; dermatological; antiallergic; protozoacide; antirheumatic; antiarthritic; osteopathic; vasotropic; analgesic; antidiabetic; nephrotropic; antianaemic; nootropic; anticonvulsant; dermatological; antigout; antiparkinsonian; antidiab
                                                                                                                                                                                                                                                                                               antibody kappa chain variable
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                                                                                                                                                               RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK
                                                                                                                                                                          Huang
                                                                                 21
                                                                                                     Indels
                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Martin F,
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                                                                                 , DB 7;
5.8e-36;
                                                                                Score 541; DB Pred. No. 5.8e 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ly that specifically for treating IL-1 nosteoarthritis and
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                                                                                 95.2%;
                                                                                                                                                                                                                                                                                              Anti-interleukin-1 receptor
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                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antibody
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                                                                                            Similarity
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                                                                                                    103;
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G;
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C IL-1 mediated diseases or for detecting the amount of IL-1R1 in a sample.

C IL-1 mediated diseases include acute pancreatitis, amyotrophic lateral sclerosis, Alzheimer's disease, cachexia, anorexia, asthma, atherosclerosis, autoimmune vasculitis, chronic fatigue syndrome, clostridium associated illnesses, coronary conditions, cancer including leukaemia and tumour metastasis, diabetes, endometriosis, fever, fibromyalgia, glomerulonephritis, graft versus host disease, costeoarthritis, rheumatoid arthritis, inflammatory eye disease, ischaemia, Kawasaki's disease, learning impairment, lung disease, ischaemia, kawasaki's disease, learning impairment, lung disease, costeocottal disease, pre-term labour, psoriasis, reperfusion injury, septic shock, side effects of radiation therapy, temporal mandibular joint disease, sleep disturbance, uveitis, or an inflammatory condition cresulting from strain, sprain, cartilage damage, trauma, orthopaedic surgery, infection or other disease processes.
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1, e.g. cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BIVLTOSPATLSLSPGERATLSCRASOSVSSYLAWYOOKPGOAPRLLIYDASNRATGIPA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 128;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human anti-TNF antibody 9E7 light chain variable region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 540; DB 8;
Pred. No. 4.3e-36;
Mismatches
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44. .54
/note= "CDR1"
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.118
.= "CDR3"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                95.1%;
96.3%;
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109. 1
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Best Local Similarity
Matches 104; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 128 AA;
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The present sequence is that of the light chain variable region of human anti-tumour necrosis factor (TNF) monoclonal antibody 9E7. This human TNF reactive IgG monoclonal antibody was generated by cloning variable and constant region DNA in vector pC4 and expression in CHO cells. The invention provides isolated human, primate, rodent, mammalian, chimeric, humanized and/or CDR-grafted anti-TNF antibodies, immunoglobulins, their cleavage products, other specified portions and variants, as well as anti-TNF antibody compositions, nucleic acids encoding these, vectors, host cells, methods for producing the antibodies using a host cell, transgenic animal or transgenic plant or plant cell, and therapeutic compositions, methods and devices. The antibody, nucleic acid, protein, composition and methods are useful for diagnosing or treating an anti-TNF related condition, e.g. cancer, anorexia, cachexia, or an immune, cardiovascular, infectious, and/or neurological disease.
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                                                                                                                                                                                      and
                                                                                                                                                           transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tumour necrosis factor; TNF; immunotherapy; TNF related diseases; obesity; immune related disease; rheumatoid arthritis; cardiovascular disease; stroke; malignant disease; leukaemia; neurological disease; multiple sclerosis; infection; hepatitis; anorectic; antiarthritic; cerebroprotective; vasotropic; cytostatic; neuroprotective; antibacterial; antiinflammatory; hepatotropic; virucide;
                                                                                                                               as anti
                                                                                                                                                                                                                 cardiovascular,
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                                                                                                                                                                                                                                                                                        Length 130;
                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                      Score 540; DB 7;
Pred. No. 4.3e-36;
; Mismatches 3
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              English
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96.3%;
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                                                                                                                                                                                                                                                             Sequence 130 AA;
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human; 9E7LC;
              Fig
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antibody

Novel isolated mammalian anti-tumor necrosis factor (TNF) alpha antibox capable of inhibiting binding of TNF alpha to TNF receptor, useful for treating TNF-related diseases such as obesity or rheumatoid arthritis.

WPI; 2004-676151/66 N-PSDB; ADS64661.

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The present invention relates to a mammalian anti-tumour necrosis factor (TNF) antibody capable of inhibiting binding of TNF alpha to TNF receptor. The invention is useful for diagnosing or treating an anti-TNF related condition in a cell, tissue, organ or animal and in immunotherapy. The invention is also useful for treating TNF related diseases chosen from obesity, immune related disease such as rheumatoid arthritis, cardiovascular disease such as stroke, malignant disease such as leukaemia, neurological disease such as multiple sclerosis and bacterial or viral infections such as hepatitis. The present sequence is the human 9E7LC light chain protein (variable region and J-region). This sequence is used in the exemplification of the invention.
                              Example 4; Fig 3; 45pp; English
xxxyyyyyyxx
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Sequence 130 AA;

ó; Gaps ô 130; Indels Length Score 540; DB 8; Le Pred. No. 4.3e-36; 1; 95.1**%**; 96.3**%**; Query Match Best Local Similarity 96.3 Matches 104; Conservative ò

108 RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK **61** ઠે 8

a a

128 81

Search completed: December 28, 2005, 13:58:08 Job time: 128.462 secs

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GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd
                 Copyright
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sw model using protein search, OM protein 13:46:34 2005, 28, December Run on:

; Search time 22.1538 Seconds (without alignments) 469.057 Million cell updates/sec

Title:

× ....QQRSNWPPFTFGPGTKVDI RIVLTQSPATLSLSPGERAT. US-09-920-137F-8 568 1 EIVLTQSPATLSLS1 score: Sequence: Perfect

108

Gapext Gapop 10.0 **BLOSUM62** Scoring table:

residues 283416 seqs, 96216763 Searched:

0

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283416 of hits satisfying chosen parameters: Total number

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

\*:08 PIR Database

pirl:\* pir2:\* pir3:\* H 20 E 4

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Result	ç	Query	ָרָ פָּיָרָ לַלְ	ä	SUMMARIES	,, ,, ,,
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### **ALIGNMENTS**

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PIDN: CAA58108.1;
                                                                                                                                                                                                                                            of
                                                                                                                                                                                                                                             domains
RESULT 1
S54905
Ig kappa chain V region - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 08-Jul-1995 #sequence_revision 21-Jul-1995 #text_change 21-Jan-20
C;Accession: S54905
R;Esposito, G.; Traboni, C.
R;Esposito, G.; Traboni, C.
R;Espositorion: Cloning and sequencing of cDNA coding for the variable dom A;Beference number: S54905
A;Accession: S54905
A;Accession: S54905
A;Residues: 1-114 <ESP>
A;Residues: 1-114 <ESP>
A;Cross-references: UNIPARC:UPI000011620A; EMBL:X82934; NID:g809554; PIDN C;Superfamily: immunoglobulin V region; immunoglobulin
F;16-90/Domain: immunoglobulin homology <IMM>
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Length 114; 7

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1 EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA Gaps ő Indels 4, Score 528; DB 2 Pred. No. 5e-39; 4; Mismatches 4. 93.0%; llarity 92.6%; Conservative Local Similarity 100; Query Match Best Local S Matches 100

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Length 111; 5 DB 524.5; Score 92.38; Query Match

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nes 100; Conservative
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C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 19-Oct-1995 #sequence_revision 19-Oct-1995 #text_change 21-Ja
C; Accession: A56701
R; Nickerson, K.G.; Tao, M.H.; Chen, H.T.; Larrick, J.; Kabat, E.A.
J. Biol. Chem. 270, 12457-12465, 1995
A; Title: Human and mouse monoclonal antibodies to blood group A subst A; Reference number: A56701; MUID:95279371; PMID:7759488
A; Accession: A56701
A; Status: preliminary
A; Molecule type: mRNA
A; Residues: 1-128 < NIC>
A; Cross-references: UNIPARC:UPI0000113B26; GB:L41174; NID:9762823; PI
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
                     1;
                                                                                                                                                                                                                                                              Ign kappa chain precursor V-J-C region (LS1) - human (fragment)
C; Species: Homo sapiens (man)
C; Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 21-c; Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 21-c; Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 21-c; Accession: PL0106
R; Silberstein, L.E.; Litwin, S.; Carmack, C.E.
J. Exp. Med. 169, 1631-1643, 1989
A; Title: Relationship of variable region genes expressed by a human A; Reference number: PL0106; MUID:89235583; PMID:2541221
A; Reference number: PL0106; MUID:89235583; PMID:2541221
A; Residues: 1-144 <SIL.>
A; Cross-references: UNIPARC; UP100001767A2
C; Superfamily: immunoglobulin V region; immunoglobulin
F; 1-20/Domain: signal sequence #status predicted <SIG>C; Keywords: heterotetramer; immunoglobulin
F; 1-20/Domain: signal sequence #status predicted <SIG>F; 1-10/Domain: immunoglobulin homology <IMM>F; 1-20/Domain: complementarity-determining 2
F; 70-76/Region: complementarity-determining 3
F; 109-115/Region: complementarity-determining 3
F; 116-127/Domain: J region <JRG>F; 118-144/Domain: C region (fragment) <CRE>F; 118-144/Domain: C region (fragment)
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                                                                                                                                  RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK
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                     Indels
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                                                           EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 519.5; DB 2;
Pred. No. 3.4e-38;
2; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 517.5; DB 2, Pred. No. 4.5e-38; 2; Mismatches 3,
   9.8e-39
        ed. No. 9.80
Mismatches
  No
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   Pred.
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Local Similarity 94.4%;
nes 102; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     / Match
Local Similarity 94.4%;
nes 102; Conservative
95.4%;
                      Conservative
 Similarity
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                 103;
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Best Local S
Matches 102
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Best Local S
Matches 102
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Grapa chain V region (JM-10) - human (fragment)
C; Species: Homo sapiens (man)
C; Species: Homo sapiens (man)
C; Date: 27-Jun-1994 #sequence_revision 27-Jun-1994 #text_change 21-Jan-2000
C; Accession: G44151
R; Zebedee, S.L; Barbas III, C.F.; Hom, Y.L.; Caothien, R.H.; Graff, R.; DeGraw, J.; Pya Proc. Natl. Acad. Sci. U.S.A. 89, 3175-3179, 1992
A; Title: Human combinatorial antibody libraries to hepatitis B surface antigen.
A; Reference number: A44151; MUID:9228746; PMID:1373487
A; Reference number: A44151
A; Reference number: A44151
A; Residues: 1-108 compared with conceptual translation
A; Molecule type: mRNA
A; Residues: 1-108 czEB>
A; Cross-references: UNIPARC:UPI00001139AC; GB:M88317; NID:g183968; PIDN:AAA35975.1; PID: A; Note: nucleotide translation not given
C; Superfamily: immunoglobulin V region; immunoglobulin
C; Superfamily: immunoglobulin homology < IMM>
F;15-89/Domain: immunoglobulin homology < IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PIDN: CAA51123.1;
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                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            hypermutation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NID:9441378;
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                                                                                                                                                                                                                                                                                  Ig kappa chain V-J-C region - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 19-May-1994 #sequence_revision 26-May-1995 #text_change 21-C;Accession: $40345
R;Klein, R.; Jaenichen, R.; Zachau, H.G.
Eur. J. Immunol. 23, 3248-3271, 1993
A;Title: Expressed human immunoglobulin chi genes and their hypermu A;Title: Expressed human immunoglobulin chi genes and their hypermu A;Reference number: $40312; MUID: 94080891; PMID: 8258341
A;Reference number: $40345
A;Katus: preliminary; translation not shown
A;Molecule type: mRNA
A;Kesidues: 1-128 <KLE>
A;Cross-references: UNIPARC: UPI000011615F; EMBL: X72455; NID: 9441376
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;32-106/Domain: immunoglobulin homology <IMM>
                                                                                                         108
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                                                                                                   RFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK
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Pred. No. 1e-37;
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llarity 91.7%;
Conservative
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PIDN:CAA61149

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Gape

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C;Species: Homo sapiens (man)
C;Date: 17-Mar-1987 #sequence_revision 17-Mar-1987 #text_change 09-Jul-2004
C;Date: 17-Mar-1987 #sequence_revision 17-Mar-1987 #text_change 09-Jul-2004
C;Accession: A01900
R;Pech, M.; Zachau, H.G.
Nucleic Acids Res. 12, 9229-9236, 1984
A;Title: Immunoglobulin genes of different subgroups are interdigitated within the V-K l A;Reference number: A93549; MUID:85087932; PMID:6440122
A;Reference number: A01900
A;Molecule type: DNA
A;Residues: 1-115 <PEC>
A;Residues: 1-115 <PEC>
A;Residues: 1-115 <PEC>
A;Cross-references: UNIPROT:P04433; UNIPARC:UPI0000116D05; GB:X01668; GB:X02768; NID:933
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-Jan-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chi genes and their hypermutation.
C;Date: 10-Oct-1995 #sequence_revision i/-Nov-1372 #tcar_comments of Accession: S57444

R;Paterson, G.; Wilson, G.; Kennedy, P.G.E.; Willison, H.J.

submitted to the EMBL Data Library, June 1995

A;Description: Analysis of anti-GM1 ganglioside IgM antibodies cloned from the analysis of anti-GM1 ganglioside IgM antibodies cloned from the approximation: S57444

A;Reference number: S57408

A;Reference number: S57408

A;Reference number: S57408

A;Reference number: S57408

A;Residues: 1-107 <PAT>
A;Residues: 1-107 <PAT>
A;Residues: 1-107 <PAT>
A;Residues: I-107 <PAT>
A;Residues: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ig kappa chain V-J region - human

C;Species: Homo sapiens (man)
C;Date: 19-May-1994 #sequence_revision 26-May-1995 #text_change 21-Jan-2(C;Accession: $40344
C;Accession: $40344
R;Klein, R.; Jaenichen, R.; Zachau, H.G.
Eur. J. Immunol. 23, 3248-3271, 1993
A;Title: Expressed human immunoglobulin chi genes and their hypermutation
A;Reference number: $40312; MUID:94080891; PMID:8258341
A;Reference number: $40312; MUID:94080891; PMID:8258341
A;Reference number: $40314
A;Reference number: $40314
A;Residues: preliminary; translation not shown
A;Molecule type: mRNA
A;Residues: 1-125 <KLE>
A;Cross-references: UNIPARC:UPI000011615E; EMBL:X72454; NID:9441376; PID?
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;36-110/Domain: immunoglobulin homology <IMM>
                                                                                                                                         cloned
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Species: Homo sapiens (man)
Date: 17-Mar-1987 #sequence_revision 17-Mar-1987 #text_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 494.5; DB 2;
Pred. No. 3.7e-36;
1; Mismatches 7;
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Pred. No. 1.4e-35;
3; Mismatches 6;
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89.8%;
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Best Local Similarity 90.6%;
Matches 96; Conservative
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Best Local Similarity
Matches 97; Conser
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S40344
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                                                             Ig kappa chain V region (60.3 hybridoma) - human C; Species: Homo sapiens (man) C; Species: Homo sapiens (man) C; Species: Homo sapiens (man) C; Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 21-Jan-2000 C; Accession: S34110; S29627 R; Walls, M.A.; Hsiao, K.; Harris, L.J. R; Walls, M.A.; Hsiao, K.; Harris, L.J. Nucleic Acids Res. 21, 2921-2929, 1993 A; Title: Vectors for the expression of PCR-amplified immunoglobulin variable domains with A; Reference number: S34110; MUID: 93324379; PMID: 9332501 A; Reference number: S34110 A; Rote: Human sequence not shown; translation not shown A; Rote: human sequences cloned and sequenced prior to expression in mouse myeloma cells A; Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1992 C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PI
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C; Species: Homo sapiens (man)
C; Date: 19-May-1994 #sequence_revision 26-May-1995 #text_change 21-Jan-C; Date: 19-May-1994 #sequence_revision 26-May-1995 #text_change 21-Jan-C; Accession: S40379
R; Klein, R.; Jaenichen, R.; Zachau, H.G.
Eur. J. Immunol. 23, 3248-3271, 1993
A; Title: Expressed human immunoglobulin chi genes and their hypermutation A; Reference number: S40312; MUID:94080891; PMID:8258341
A; Accession: S40379
A; Accession: S40379
A; Catus: preliminary; translation not shown
A; Molecule type: mRNA
A; Residues: 1-128 <KLE>
A; Cross-references: UNIPARC:UPI000116181; EMBL:X72489; NID:9441446; PII
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F; 36-110/Domain: immunoglobulin homology <IMM>
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C;Superfamily: immunoglobulin V region; immunoglobulin
C;Keywords: heterotetramer; immunoglobulin
F;36-110/Domain: immunoglobulin homology <IMM>
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Pred. No. 1.8e-37;
1; Mismatches 6;
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           A; Note: the Sequence was determined from the germine general C; Genetics:
A; Genetics:
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A; Complex: An immunoglobulin heterotetramer subunit consists of two ident c; Complex: An immunoglobulin beterotetramer; immunoglobulin homology
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Reywords: heterotetramer; immunoglobulin predicted <SIG>F; 1-20 / Domain: signal sequence #status predicted <SIG>F; 21-115 / Product: Ig kappa chain V-III region (Vg) #status predicted <MA; F; 21-43 / Region: framework 1
F; 36-110 / Domain: immunoglobulin homology <IMM>F; 36-110 / Domain: framework 2
F; 70-76 / Region: complementarity-determining 2
F; 70-76 / Region: framework 3
F; 109-115 / Region: complementarity-determining 3
F; 109-115 / Region: complementarity-determining 3
F; 109-115 / Region: complementarity-determining 3
F; 109-115 / Region: status predicted
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C; Species: Homo sapiens (man)
C; Date: 05-Dec-1998 #sequence_revision 05-Dec-1998 #text_change 21-Jan-C; Accession: JE0244
R; Alim, M.A.; Hara, Y.; Hossain, M.S.; Takeda, K.; Yamagata, F.; Yamaki submitted to JIPID, November 1998
A; Description: A new subgroup of k type light chains (VkV) identified i A; Reference number: JE0243
A; Accession: JE0244
A; Molecule type: protein
A; Residues: 1-215 <ALI>A; Residues: 1-215 <ALI>A; Cross-references: UNIPARC:UPI0000176982
C; Superfamily: immunoglobulin V region; immunoglobulin homology
F; 16-90/Domain: immunoglobulin homology <IMM>
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C; Species: 07-Jul-1997 #sequence_revision 29-Aug-1997 #text_change 21-C; Date: 07-Jul-1997 #sequence_revision 29-Aug-1997 #text_change 21-C; Accession: PC4282; PC4284
R; Suzuki, H.; Takemura, H.; Suzuki, M.; Sekine, Y.; Kashiwagi, H. Biochem. Biophys. Res. Commun. 232, 101-106, 1997
A; Title: Molecular cloning of anti-ss-A/Ro 60-kDa peptide fab fragm
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Mismatches
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Pred. No. 1.8e-35
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C; Species: Homo sapiens (man)
C; Date: 25-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 21-Jan-20 C; Accession: S11697
R; Pech, M.; Smola, H.; Pohlenz, H.D.; Straubinger, B.; Gerl, R.; Zachau, J. Mol. Biol. 183, 291-299, 1985
A; Title: A large section of the gene locus encoding human immunoglobulin A; Reference number: S11697; MUID: 85264787; PMID: 3927006
A; Accession: S11697
A; Status: preliminary; nucleic acid sequence not shown; translation not shimolecule type: DNA
A; Residues: 1-115 < PEC>
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C;Date: 27-Jan-1995 #sequence_revision 01-Sep-1995 #text_change 21-Jan-20;Date: 27-Jan-1995 #sequence_revision 01-Sep-1995 #text_change 21-Jan-20;Accession: $46375; $38648
R;Bensimon, C.; Chastagner, P.; Zouali, M.
EMBO J. 13, 2951-2962, 1994
A;Title: Human lupus anti-DNA autoantibodies undergo essentially primary A;Reference number: $46369; MUID:94313975; PMID:8039491
A;Reference number: $46375
A;Molecule type: mRNA
A;Residues: 1-114 <BEN>
A;Residues: 1-114 <BEN>
Cross-references: UNIPARC:UPI00001165A8; EMBL:Z27176; NID:g415967; PIDIC; Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;20-95/Domain: immunoglobulin homology <IMM>
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Pred. No. 3.2e-34;
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                 A; Accession: PC4282
A; Molecule type: protein
A; Residues: 1-106 <SUZ>
A; Note: E-42
A; Note: E-42
A; Accession: PC4284
A; Molecule type: protein
A; Residues: 1-106 <SUZ>
A; Cross-references: UNIPARC: UPI0000176E92
A; Cross-references: UNIPARC: UPI0000176E92
A; Note: E-56
C; Comment: This antibody is commonly found if C; Superfamily: immunoglobulin V region; immu F; 14-88/Domain: immunoglobulin homology <IMP
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A;Cross-references: UNIPARC:UPI0000115E41; EMBL:X17264; NID:g37898; PIDN:CAA35168.1; PID A;Note: the nucleotide sequence was submitted to the EMBL Data Library, December 1989 C;Genetics:
A;Introns: 17/1
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;36-110/Domain: immunoglobulin homology <IMM>
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Gaps ö Length 115; Indels Score 471; DB 2; I Pred. No. 4.3e-34; 0; Mismatches 3; Query Match
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2005, 13:44:24 ; Search time 137.077 Seconds (without alignments) 555.871 Million cell updates/sec December 28, Run on:

US-09-920-137F-8 568 1 EIVLTQSPATLSLSPGERAT......QQRSNWPPFTFGPGTKVDIK 108 Title: Perfect score: Seguence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

2166443 seqs, 705528306 residues Searched:

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

UniProt\_05.80:\*
1: uniprot\_sprot:\*
2: uniprot\_trembl:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

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"V kappa gene segments rearranged in chronic lymphocytic leukemia distributed over a large portion of the V kappa locus and do not segmentic mutation.";
Eur. J. Immunol. 23:391-397(1993).
EMBL; AF035031; AAD56267.1; -; mRNA.
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NUCLEOTIDE SEQUENCE.
MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;
Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
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UL83 HUMAN
Q9UL83 HUMAN PRELIMINARY; PRT; 108 AA.
Q9UL83;
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 25, Last annotation update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Myosin-reactive immunoglobulin light chain variable reg
(Fragment).
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Pred. No. 5.3e-42;
5; Mismatches 7;
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                                                                                                                129
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EMBL; AF035031; AAD56267.1; -; mRNAPIR; B30609; B30609.

PIR; C30609; C30609.

PIR; D30609; D30609.

PIR; S34098; S34098.

PIR; S34099; S34099.

HSSP; P01625; 1LVE.

SMR; Q9UL83; 1-108.
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illarity 88.0%;
Conservative
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95; Conserv
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108 AA;
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ID KV3H HUMAN
AC P04207;
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Best Local S
Matches 95
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                                                                                                                                                                                                                                                                                                         This Swiss-Prot entry is copyright. It is produced through a collaboratic between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use as long as its content is in no way modified and this statement is no
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
20-MAR-1987 (Rel. 04, Created)
01-NOV-1990 (Rel. 16, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Ig kappa chain V-III region CLL precursor (Rheumatoid factor).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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Complementarity-determining-1
Framework-2.
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Complementarity-determining-
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Immunoglobulin domain; Immunoglobulin V region; Signal
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Q9UL85;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Myosin-reactive immunoglobulin kappa chain variable
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Pred. No. 2.5e-41
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Mismatches
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                                                                                                                                                                      NUCLEOTIDE SEQUENCE.
MEDLINE=86177570; PubMed=3083417;
Jirik F.R., Sorge J., Fong S., Heitzmann Goldfien R., Carson D.A.;
"Cloning and sequence determination of a light-chain gene.";
Proc. Natl. Acad. Sci. U.S.A. 83:2195-219
                                                                                                                                                                                                                                                                                                                                                                                                                 Genomic_DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                             GO; GO: 0005576; C: extracellular region; GO; GO: 0003823; F: antigen binding; NAS. GO; GO: 0006955; P: immune response; NAS. InterPro; IPR007110; Ig-like. InterPro; IPR003596; Ig_v. SMART; SM00406; IGV; 1.
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ilarity 83.3%;
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HSSP; P01625; 1LVE.
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NCBI_TaxID=9606;
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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DISEASE: The protein is one of the surface immunoglobulin M autoantibodies expressed in patients with chronic lymphocytic
                                                                                                                                                                                                                                                    and norm
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                                                                                                                                      MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531 Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUCLEOTIDE SEQUENCE.
MEDLINE=88171307; PubMed=3127527; DOI=10.1084/jem.167.3.840;
Kipps T.J., Tomhave E., Chen P.P., Carson D.A.;
"Autoantibody-associated kappa light chain variable region gent expressed in chronic lymphocytic leukemia with little or no mutation. Implications for etiology and immunotherapy.";
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Clin. Immunol. Immunopathol. 87:184-192(1998).
Clin. Immunol. Immunopathol. 87:184-192(1998).
EMBL; AF035029; AAD56265.1; -; mRNA.
PIR; D30609; D30609.
HSSP; P01625; 1EK3.
SMR; Q9UL85; 1-109.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
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8.5e-41;
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01-NOV-1990 (Rel. 16, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Ig kappa chain V-III region HAH precursor.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Verte
                                                                                                                                                                                                                                                    autoantibodies in rheumatic
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HSSP; P01625; 1EEQ.
SMR; P18135; 21-129.
Ensembl; ENSG0000169769; Homo sapiens.
GO; GO:0005576; C:extracellular region;
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 468;
Pred. No.
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82.4%;
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                                                                                                            SEQUENCE
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                                                                                                                                                                                                                                                    "Myosin-reactive
                                      TaxID=9606;
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Young D.C.;
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                                                                                                            NUCLEOTIDE
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HUMAN
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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
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21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Ig kappa chain V-III region POM.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PROTEIN SEQUENCE.

MEDLINE=76276460; PubMed=60899;

Klapper D.G., Capra J.D.;

"The amino acid sequence of the variable regions of the light chains from two idiotypically cross reactive IgM anti-gamma globulins.";

Ann. Immunol. (Paris) 127C:261-271(1976).

-!- MISCELLANEOUS: This chain was isolated from an IgM with anti-gamma
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Complementarity-determining-3
JK1 segment.
By similarity.
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Immunoglobulin domain;
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SMR; P01624; 1-109.
SMR; P01624; 1-109.
GO; GO: 0005576; C: extracellular region; NA
GO; GO: 0003823; F: antigen binding; NAS.
GO; GO: 0006955; P: immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig-v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
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InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Immunoglobulin domain; Immunog
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Immunoglobulin V region.
DISULFID 23 89
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Wu X., Liu
Young D.C.;
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-!- DISEASE: The protein is one of the surface immunoglobulin autoantibodies expressed in patients with chronic lymphocy
                                                                                  2;
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Framework-1.
Complementarity-determining-1
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                 62821DDC6A8ABA86 CRC64;
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                                                 Score 453; DB 1;
Pred. No. 3.1e-39
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01-NOV-1990 (Rel. 16, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Ig kappa chain V-III region HIC precursor.
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InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Immunoglobulin domain; Immunoglobulin V response.
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P; P01625; 1EEQ.
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embl; ENSG00000169769; 1
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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                109 AA;
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10-MAY-2005
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SMR; P18136;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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A Martin T., Duffy S.F., Carson D.A., Kipps T.J.;

Martin T., Duffy S.F., Carson D.A., Kipps T.J.;

"Evidence for somatic selection of natural autoantibodies.";

EMBL; A30601; A30601.

PIR; A30601; B30601.

PIR; C30601; B30601.

PIR; D30607; D30607.

PIR; D30607; D30607.

PIR; F30608; F30608.

PIR; F30608; F30608.

PIR; G30608; G30608.

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PubMed=1373487;
Zebedee S.L., Barbas C.F. 3rd, Hom Y.L., Caothien R.H., Graff I
DeGraw J., Pyati J., LaPolla R., Burton D.R., Lerner R.A.;
"Human combinatorial antibody libraries to hepatitis B surface
                       2;
                                                                                                                                                                                                        128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531 Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                     ARFSGSGSGTDFTLTISSLEPEDFAVYYCQORSNWPPFTFGPGTKVDIK
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin light chain variable
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 Pred. No. 3.8e-39; Mismatches 1
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Clin. Immunol. Immunopathol. 87:184-192(1998)
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"V kappa gene segments rearranged in distributed over a large portion of t somatic mutation.";
                                                                                                                                                                                                                                                                                                                        PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Myosin-reactive autoantibodies in
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85.3%;
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Q9UL78 HUMAN PRELIMINARY
Q9UL78;
 Similarity 85... 93; Conservative
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PubMed=8436174;
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Proc. Natl. Acad.
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KV3D HUMAN
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MEDLINE=82046598; PubMed=6794615;
Andrews D.W., Capra J.D.;
Andrews D.W., Capra J.D.;
"Amino acid sequence of the variable regions of light chains from two idiotypically cross-reactive human IgM anti-gamma-globulins of the Wa group.";
Biochemistry 20:5816-5822(1981).
Biochemistry 20:5816-5822(1981).
-!- MISCELLANEOUS: This chain was isolated from an IgM with anti-gamma globulin activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Ig kappa chain V-III region SIE.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                         BIVLTQSPATLSLSPGERATLSCRASQSV-YSYLAWYQQKPGQAPRLLIYDASNRATGIP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRSNWPPFTFGPGTKVDIK 108
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Immunoglobulin domain;
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Pred. No. 8.1e-39;
; Mismatches 11
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HSSP; P01625; 1LVE.

SMR; P01620; 1-109.

GO; GO:0003823; F:antigen binding; NAS.

GO; GO:0006955; P:immune response; NAS.

InterPro; IPR007110; Ig-like.

InterPro; IPR003596; Ig_v.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIKE; 1.

Direct protein sequencing; Immunoglobulin doma Immunoglobulin V region.

DISULFID 23 89 By similarity.
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nilarity 85.3%;
Conservative
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InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
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                                        PH0963.
PH0964.
PH0965.
S33988.
S34096.
PIR; H44151; H44151
PIR; I30601; I30601
PIR; PH0963; PH0963
PIR; PH0964; PH0964
PIR; PH0965; PH0965
PIR; S33988; S33988
PIR; S34096; S34096
HSSP; P01625; 1EK3.
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93; Conserv
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PROSITE; PS50835;
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21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Ig kappa chain V-III region Ti.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Bence-Jones protein; Direct protein sequencing; Immunoglobulin V region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Suter L., Barnikol H.U., Watanabe S., Hilschmann N.;
"Rule of antibody structure. The primary structure of a monoclonal immunoglobulin L-chain of kappa-type, subgroup 3 (Bence-Jones protein Ti). IV. The complete amino acid sequence and its significance for the mechanism of antibody production.";
Hoppe-Seyler's Z. Physiol. Chem. 353:189-208(1972).
-!- MISCELLANEOUS: The C region of this chain has the INV (3) marker.
                                                                                                                             1 BIVLTQSPATLSLSPGERATLSCRASQSV-YSYLAWYQQKPGQAPRLLIYDASNRATGIP
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                CRC64;
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Pred. No. 1e-38;
; Mismatches
                                                                      1e-38;
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HSSP; P01625; 1LVE.
SMR; P01622; 1-109.
GO; GO:0005576; C:extracellular region; NA
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
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84.4%;
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92; Conserv
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Hypothetical protein.
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TISSUE=Brain;
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05-JUL-2004
05-JUL-2004
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Best Local S
Matches 90
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Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                  Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                     in rheumatic carditis and norm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Brain;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler
Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.
                                                                                                                                                                          MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531 Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                              109;
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                                                                       01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
Myosin-reactive immunoglobulin kappa chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                12;
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                                                                                                                                                                                                                  87:184-192(1998).
-; mRNA.
                                                                                                                                                                                                                                                                                                                                                             Score 448; DB 2
Pred. No. 1e-38;
2; Mismatches
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HSSP; P01625; IEK3.
SMR; Q9UL86; 1-109.
Ensembl; ENSG00000169769; Homo sapiens.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
InterPro; IPR003596; Ig_v.
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85.3%;
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Q9UL86_HUMAN PRELIMINARY;
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Homo sapiens (Human).
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PIR; B30607; B30607.
PIR; I30601; I30601.
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                                                                                                            Homo sapiens (Human)
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Mammalia; Eutheria;
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93; Conserv
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109 AA;
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05-JUL-2004
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Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
Generation and initial analysis of more than 15,000 full-length human
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Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
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Strausberg R.;
Submitted (JUN-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; BC032451; AAH32451.1; -; mRNA.
HSSP; P01837; 1KCU.
SMR; Q6PILB; 21-236.
InterPro; IPR003599; Ig.
InterPro; IPR003599; Ig.
InterPro; IPR003599; Ig.
InterPro; IPR003596; Ig.
InterPro; IPR003596; Ig.
InterPro; IPR003596; Ig.
SMART; SM00409; IG; 2.
SMART; SM00400; IGc1; 1.
SMART; SM00406; IGv; 1.
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Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                           and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002)
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Pred. No. 2.9e-38;
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PROSITE; PS00290; IG MHC; UNKNOWN 1.
Hypothetical protein.
SEQUENCE 236 AA; 25834 MW.
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82.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A., Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.; "Generation analysis of more than 15,000 full-length human Generation analysis of more than 15,000 full-length human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUCLECTIDE SEQUENCE.

TISSUE=Glandular pool- thyroid;

MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler (Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.)

Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheet:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Last sequence update)
Last annotation update)
                                                                                                                                                    "Generation and sequences.";
and mouse cDNA sequences.";
..., Arad. Sci. U.S.A. 99:16899-16903(2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 447; DB 2; 1
Pred. No. 3.3e-38;
5; Mismatches 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                          F33A145A396BA285
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                                                                                                                                                                                                                                                                                                                                                                SMART; SM00409; IG; 2.
SMART; SM00407; IGc1; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 2.
PROSITE; PS00290; IG MHC; UNKNOWN 1.
Hypothetical protein.
SEQUENCE 235 AA; 25520 MW; F33A14
                                                                                                                                                                                                                                                     -; mRNA.
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N
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Strausberg R.;
Submitted (OCT-2001) to the E
EMBL; BC016380; AAH16380.1; -
HSSP; P01837; 1KCU.
SMR; Q6PJF2; 21-235.
InterPro; IPR003599; Ig.
InterPro; IPR003597; Ig-like.
InterPro; IPR003596; Ig-MHC.
InterPro; IPR003596; Ig_V.
                                                                                                                                                                                                                                                                                                 19-like.
19-cl.
19-MHC.
19-v.
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27,
27,
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05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
Hypothetical protein.
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Bukaryota; Metazoa;
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Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McBwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
Chnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
The Generation and initial analysis of more than 15,000 full-length human
The Proc. Natl. Acad. Sci. U.S.A. 99:16899-16007/2007
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21-JUL-1986 (Rel. 01, Last sequence update)
10-MAY-2005 (Rel. 47, Last annotation update)
Ig kappa chain V-III region WOL.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
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SRATGIP
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Andrews D.W., Capra J.D.;
"Amino acid sequence of the variable regions of light chains
idiotypically cross-reactive human IgM anti-gamma-globulins o
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NO. 7.6e-38;
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Pred. No. 7.6e
9; Mismatches
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WEOUS: This chain was isolated
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PROSITE; PS00290; IG MHC; UNKNOWN_1.
Hypothetical protein.
SEQUENCE 236 AA; 25773 MW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -; mRNA
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SMR; Q6P5S8; 21-236.
InterPro; IPR003599; IG.
InterPro; IPR003109; Ig_cl.
InterPro; IPR003597; Ig_cl.
InterPro; IPR00306; Ig_MHC.
InterPro; IPR00306; Ig_WHC.
InterPro; IPR00306; Ig_W.
Pfam; PF07654; Cl-set; 1.
SMART; SM00409; IG; 2.
SMART; SM00407; IGC1; 1.
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illarity 79.8%;
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Biochemistry 20:5816-582
-!- MISCELLANEOUS: This
globulin activity.
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C
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Submitted (NOV-2003)
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87; Conser
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PROTEIN SEQUENCE.
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This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use as long as its content is in no way modified and this statement is not
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HSSP; P01625; 1LVE.
SMR; P01623; 1-109.
GO; GO:0005576; C:extracellular region; NAS.
GO; GO:0006955; P:immune response; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR00310; Ig-like.
InterPro; IPR003596; Ig-v.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 1.
Direct protein sequencing; Immunoglobulin domain; Immunoglobulin V region.
DISULFID 23 89 By similarity.
NON TER 109 AA; 11746 MW; 566C115E6B9CBEEE CRC6.
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Pred. No. 4.3e-38;
3; Mismatches 12;
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Search completed: December 28, 2005, 14:03:09 Job time : 138.077 secs

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                                                                APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
FILE REFERENCE: 020015-000200US
CURRENT APPLICATION NUMBER: US/09/456,090A
CURRENT FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 110
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 46
LENGTH: 224
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Pred. No. 2e-42;
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            Sequence 46, Application US/09456090A Patent No. 6680209 GENERAL INFORMATION:
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Best Local Similarity 93.5%;
Matches 101; Conservative
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ORGANISM: Homo sapiens
OTHER INFORMATION: M1-10L
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APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: GenPharm International
TITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 020015-000110US
CURRENT APPLICATION NUMBER: US/09/453,234

Sequence 46, Application US/09453234 Sequence 46, Application US/09453234 Patent No. 6794132 GENERAL INFORMATION: APPLICANT: Buechler, Joe

RESULT US-09-4

118

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TYPE: PRT
ORGANISM: Homo sapiens
OTHER INFORMATION: M2-31L
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ORGANISM: Homo sapiens
OTHER INFORMATION: M2-34L
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SEQ ID NO 82
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Sequence 2, Application US/09434870
Sequence 2, Application US/09434870
GENERAL INFORMATION:
APPLICANT: Huse, William
APPLICANT: Wu, Herren
TITLE OF INVENTION: Methods of Optimizing Antibody Variable Region Binding Affinity
FILE REFERENCE: AME-06352
CURRENT APPLICATION NUMBER: US/09/434,870
CURRENT FILING DATE: 2001-08-20
PRIOR PPLICATION NUMBER: 60/159,689
PRIOR FILING DATE: 1999-10-14
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APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff; APPLICANT: Lonberg, Nils
TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGN
FILE REFERENCE: 020015-000200US
CURRENT APPLICATION NUMBER: US/09/456,090A
CURRENT FILING DATE: 1999-12-06
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Best Local Similarity 95.4%;
Matches 103; Conservative
           PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 46
LENGTH: 224
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SOFTWARE: Patentin version
                                                                                                    ORGANISM: Homo sapiens
COTHER INFORMATION: M1-10L
US-09-453-234-46
  NUMBER:
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                                                                                                                                                                  Query Match
Best Local Similarity
Matches 101; Conser
APPLICATION
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; Sequence 90, Application US/09456090A
; GENERAL INFORMATION:
; APPLICANT: Buechler, Joe
; APPLICANT: Valkirs, Gunars
; APPLICANT: Gray, Jeff
; APPLICANT: Lonberg, Nils
; TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
; TITLE OF INVENTION: HUMAN ANTIBODIES, O90A
; TITLE NEFERENCE: 020015-000200US
; CURRENT APPLICATION NUMBER: US/09/456,090A
; CURRENT FILING DATE: 1999-12-06
; NUMBER OF SEQ ID NOS: 110
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 90
; LENGTH: 224
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                                                                                                                                                                                   Sequence 88, Application US/09456090A
Patent No. 6680209
GENERAL INFORMATION:
APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
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Pred. No. 6e-42;
2; Mismatches 3;
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 Score 518.5; DB Pred. No. 6e-42; 2; Mismatches
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CURRENT APPLICATION NUMBER: US/09/456,090A
CURRENT FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 110
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 88
LENGTH: 224
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                                                                                                                                                                                                                                                                                           Sequence 82, Application US/09453234

Patent No. 6794132

GENERAL INFORMATION:

APPLICANT: Buechler, Joe

APPLICANT: Valkirs, Gunars

APPLICANT: Lonberg, Nils

APPLICANT: GenPharm International

TITLE OF INVENTION: Human Antibodies

FILE REFERENCE: 020015-000110US

CURRENT APPLICATION NUMBER: US/09/453,234

CURRENT FILING DATE: 1999-12-01

PRIOR FILING DATE: 1999-10-02

NUMBER OF SEQ ID NOS: 112

SOFTWARE: Patentin Ver. 2.1
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CURRENT FILING DATE: 1999-12-01
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Patent No. 6794132;
GENERAL INFORMATION:
APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Gray, Jeff
APPLICANT: Gonberg, Nils
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: GenPharm International
TITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 020015-000110US
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ilarity 94.4%;
Conservative
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                                                                                     Best_Local Similarity 94.4
Matches 102; Conservative
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; ORGANISM: Homo sapiens
; OTHER INFORMATION: M2-31L
US-09-453-234-82
          ORGANISM: Homo sapiens
OTHER INFORMATION: M2-35L
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ORGANISM: ...
COTHER INFORMAT:
US-09-456-090A-90
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US-09-453-234-88
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ENGTH: 224
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APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
FILE REFERENCE: 020015-000200US
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CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT FILING DATE: 1999-12-01
PRIOR APPLICATION NUMBER: US 60/157,415
PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: GenPharm International
TITLE OF INVENTION: Human Antibodies
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Patent No. 6680209
GENERAL INFORMATION:
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Sequence 90, Application US/09453234
Patent No. 6794132
GENERAL INFORMATION:
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1999-10-02
OS: 112
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94.4%;
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Best Local Similarity 94.4%;
Matches 102; Conservative
                                                                                         ; ORGANISM: Homo sapiens
; OTHER INFORMATION: M2-34L
US-09-453-234-88
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CTHER INFORMATION: M2-35L
US-09-453-234-90
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PRIOR FILING DATE: 1999-1
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 7
SEQ ID NO 88
LENGTH: 224
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US-09-456-090A-36
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ENGTH: 224
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Patent No. 6680209
GENERAL INFORMATION:
APPLICANT: Buechler, Joe
APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
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Pred. No. 7.5e-42;
; Mismatches 4;
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Pred. No. 7.5e-42;
; Mismatches 4
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APPLICANT: Gray, Jeff
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: GenPharm International
TITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 020015-000110US
CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT FILING DATE: 1999-12-01
PRIOR APPLICATION NUMBER: US 60/157,415
PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 36
LENGTH: 224
CURRENT APPLICATION NUMBER: US/09/456,090A CURRENT FILING DATE: 1999-12-06 NUMBER OF SEQ ID NOS: 110 SOFTWARE: PatentIn Ver. 2.1
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Patent No. 6794132
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nilarity 94.4%;
Conservative
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Best Local Similarity 94.4%;
Matches 102; Conservative
                                                                                                                                                         OTHER INFORMATION: M1-1L
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APPLICANT: Buechler, Joe
                                                                                                                      ORGANISM: Homo sapiens
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US-09-453-234-36
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Best Local Similarity
Matches 102; Conser
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US-09-456-090A-84
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                                                                    SEQ ID NO 36
LENGTH: 224
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Pred. No. 1.4e
2; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: GenPharm International
TITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 020015-0001100S
CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT FILING DATE: 1999-12-01
PRIOR APPLICATION NUMBER: US 60/157,415
PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 514.5;
Pred. No. 1.4
FILE REFERENCE: 020015-000200US
CURRENT APPLICATION NUMBER: US/09/456,090A
CURRENT FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 110
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 13
US-09-453-234-84
; Sequence 84, Application US/09453234
; Patent No. 6794132
; GENERAL INFORMATION:
; APPLICANT: Buechler, Joe
; APPLICANT: Valkirs, Gunars
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 40, Application US/09438954; Patent No. 6458934; GENERAL INFORMATION:
APPLICANT: HONG, Hyo Jeong; APPLICANT: PARK, Sung Sup; APPLICANT: KANG, Young Jun; APPLICANT: KANG, Young Jun; APPLICANT: KANG, Chang-Yuil
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.larity 93.5%;
Conservative 2
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Local Similarity 93.5%;
les 101; Conservative
                                                                                                              TYPE: PRT
ORGANISM: Homo sapiens
OTHER INFORMATION: M2-32L
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ORGANISM: Homo sapiens
OTHER INFORMATION: M2-32L
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US-09-456-090A-84
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SOFTWARE: Pactor of 1D NO 84
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                                                                               SEQ ID NO 84
LENGTH: 224
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Gaps
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                                                                                                                                                                                                             Artificial Sequence: Variable chain of human antibody (X82934)
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TITLE OF INVENTION: HUMANIZED ANTIBODY SPECIFIC FOR HUMAN 4-1BB

TITLE OF INVENTION: PHARMACEUTICAL COMPOSITION COMPRISING SAME
FILE REFERENCE: 1303-124P

CURRENT APPLICATION NUMBER: US/09/438,954

CURRENT APPLICATION NUMBER: US/09/438,954

CURRENT FILING DATE: 1999-11-12

NUMBER OF SEQ ID NOS: 49

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 40

LENGTH: 107
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US-09-456-090A-38
i Sequence 38, Application US/09456090A
i Patent No. 6680209
i GENERAL INFORMATION:
i APPLICANT: Buechler, Joe
i APPLICANT: Ucnberg, Nils
i APPLICANT: Lonberg, Nils
i TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
i TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
i TITLE OF INVENTION: HUMAN ANTIBODIES
i CURRENT APPLICATION NUMBER: US/09/456,090A
i CURRENT FILING DATE: 1999-12-06
i NUMBER OF SEQ ID NOS: 110
i SEQ ID NO 38
i LENGTH: 226
i TYPE: PRT
i ORGANISM: Homo BapienB
i OTHER INFORMATION: M1-3L
US-09-456-090A-38
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Pred. No. 1.9e-41;
5; Mismatches 4;
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Pred. No. 1.5e-39;
3; Mismatches 7;
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ORGANISM: Artificial Sequence
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Similarity 90.7%;
98; Conservative
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CTHER INFORMATION:
US-09-438-954-40
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US-09-920-137F-8 568 score: Title: Perfect

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Sequence:

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# SUMMARIES

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Description	Sequence	Sequence	Sequence	Sequence	Seguence	Sequence	Sequence	Sequence	Seguence	Seguence	Seguence	Sequence	Seguence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Seguence
ΔΙ	US-09-920-267C-8	-305-347	US-10-720-323-8	US-10-954-900A-8	US-11-009-731-90	US-10-408-901-24	US-10-408-901-48	US-10-656-769-12	US-10-394-471B-16	US-10-656-769-38	US-10-408-901-8	US-10-408-901-32	US-10-173-551-9	US-10-886-838-4	21-71	US-11-009-731-85	US-11-102-403-9	US-11-102-403-11	US-10-408-901-16	US-10-408-901-28	US-10-408-901-40		US-10-478-056-19	US-10-251-085B-142	US-10-737-252-142	US-10-961-567A-6	US-09-880-748-1049
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Sequence 1049, Ap Sequence 91, Appl	equence 2, App	equence 92, Ap	, Ap	equence 46, Ap	19, A	equence 25, Ap	equence 71, Ap	equence 2, App	equence 2, App	Sequence 2, Appli	equence 14, Ap	equence 21, Ap	Sequence 3, Appli		, Ap	, Ap
US-10-293-418-1049 US-10-891-658-91	-10-886-838-	US-11-009-731-92	US-09-791-153A-69	US-09-453-234-46	US-10-173-551-19	US-10-443-466A-25	1-153A-7	US-09-982-464-2	US-10-697-399-2	US-10-697-400-2	US-10-911-838-14	US-10-911-838-21	US-11-102-403-3	US-10-478-056-31	7-799-5	US-11-021-715-99
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### ALIGNMENTS

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COMPOSITINS, METHODS AND USES
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US-10-305-347A-8
; Sequence 8, Application US/10305347A
; Publication No. US20030143603A1
; GENERAL INFORMATION:
; APPLICANT: Giles-Komar, Jill
; APPLICANT: Bernie Scallon
; TITLE OF INVENTION: ANTI-TNF ANTIBODIES, COMPOSITIONS, METHODS AND USES
; FILE REFERENCE: CENSO05
; CURRENT APPLICATION NUMBER: US/10/305,347A
; CURRENT FILING DATE: 2002-11-26
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: Patentin Ver 3.0
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    Sequence 8, Application US/09920267C;
Sequence 8, Application US/09920267C;
Publication No. US20030040044A1;
GENERAL INFORMATION:
APPLICANT: Centocor, Inc.;
APPLICANT: Trikha, Mohit;
APPLICANT: Trikha, Mohit;
APPLICANT: Snyder, Linda
APPLICANT: Nakada, Marian;
TITLE OF INVENTION: ANTI-DUAL INTEGRIN ANTIBODIES, CON;
FILE REFERENCE: CEN 249
CURRENT APPLICATION NUMBER: US/09/920,267C;
CURRENT FILING DATE: 2001-08-01
PRIOR PILING DATE: 2000-08-07
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin version 3.1
SEQ ID NO 8
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Pred. No. 1.8e-41
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Mismatches
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nilarity 99.1%;
Conservative
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US-09-920-267C-8
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Best Local Similarity
Matches 107; Conser
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US-09-920-267C-8
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APPLICANT: RANGAN, Vangipuram
APPLICANT: LANE, Thomas E.
APPLICANT: KEIRSTEAD, Hans S.
APPLICANT: LIU, Michael T.
TITLE OF INVENTION: IP-10 ANTIBODIES AND THEIR USES FILE REFERENCE: MXI-312
                                                                                                                                                                                        Score 559; DB 5;
Pred. No. 1.8e-41;
); Mismatches 1
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Pred. No. 6.7e-40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 90, Application US/11009731; Publication No. US20050191293A1; GENERAL INFORMATION:
APPLICANT: DESHPANDE, Shirkant; APPLICANT: SRINIVASAN, Mohan; APPLICANT: CARDARELLI, Josephine M.; APPLICANT: WANG, Changyu, APPLICANT: WANG, Changyu, APPLICANT: RANGAN, Vangipuram; APPLICANT: RANGAN, Vangipuram; APPLICANT: KEIRSTEAD, Hans S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 126
SOFTWARE: FastSEQ for Windows Version
PRIOR FILING DATE: 2000-08-07
PRIOR APPLICATION NUMBER: 60/236,826
PRIOR FILING DATE: 2000-09-29
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver 3.1
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US-10-408-901-24
; Sequence 24, Application US/10408901
; Publication No. US20040023313A1
; GENERAL INFORMATION:
; APPLICANT: Boyle, William
; APPLICANT: Huang, Haichun
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Best Local Similarity 99.1
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                                                                                                    LENGTH: 108
TYPE: PRT
ORGANISM: Homo sapiens
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US-11-009-731-90
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LENGTH: 108
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                                                                                                                                                                                              APPLICANT: Bernie Scallon
APPLICANT: George Heavner
TITLE OF INVENTION: ANTI-TNF ANTIBODIES, COMPOSITIONS, METHODS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EIVLTQSPATLSLSPGERATLSCRASQSVYSYLAWYQQKPGQAPRLLIYDASNRATGIPA
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                                                                                                        Length 108;
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Trikha, Mohit
Snyder, Linda
Nakada, Marian
VENTION: ANTI-DUAL INTEGRIN ANTIBODIES,
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                                                                                                       Score 559; DB 4;
Pred. No. 1.8e-41
; Mismatches
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Pred. No. 1.8e-41
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Giles-Komar, Jill
APPLICANT: Trikha, Mohit
APPLICANT: Snyder, Linda
APPLICANT: Nakada, Marian
TITLE OF INVENTION: ANTI-DUAL INTEGRIN ANTIF
FILE REFERENCE: CEN 249 CIPNP
CURRENT APPLICATION NUMBER: US/10/720,323
CURRENT FILING DATE: 2003-11-24
PRIOR APPLICATION NUMBER: 60/223,363
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CURRENT APPLICATION NUMBER: US/10/954,900A
CURRENT FILING DATE: 2004-09-30
PRIOR APPLICATION NUMBER: 09/920,262
PRIOR FILING DATE: 2001-08-01
PRIOR APPLICATION NUMBER: 60/223,360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 8, Application US/10954900A
Publication No. US20050123541A1
GENERAL INFORMATION:
APPLICANT: Giles-Komar, Jill
APPLICANT: David Shealy
APPLICANT: David Knight
                                                                                                                                                                                                                                                                                                                                                                   Sequence 8, Application US/10720323 Publication No. US20040185507A1 GENERAL INFORMATION:
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nilarity 99.1%;
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Best Local Similarity 99.1%;
Matches 107; Conservative
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SOFTWARE: Patentin version
                                                                                                                                                                                                                                                                                                                                                                                                                      , Inc.
                                                      Homo sapiens
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; ORGANISM: Homo sapiens
US-10-720-323-8
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Best Local Similarity
Matches 107; Conser
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                   ; LENGTH: 108
; TYPE: PRT
; ORGANISM: Hom
US-10-305-347A-8
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APPLICANT:
APPLICANT:
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Publication No. US20040185047A1
GENERAL INFORMATION:
APPLICANT: Giles-Komar, Jill; Carton, Jill; Scallon, Bernard J.
TITLE OF INVENTION: ANTI-TNF ANTIBODIES, COMPOSITIONS, METHODS
FILE REFERENCE: CEN0202
CURRENT APPLICATION NUMBER: US/10/394,471B
CURRENT FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/367,903
PRIOR FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver 3.1
                                                                                                                                                                                                                                                                                                                                                                                               Length 128;
                                                                                                      Anti-IL-1R Monoclonal
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Pred. No. 9.8e-40;
; Mismatches 3;
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Pred. No. 9.6e-40;
; Mismatches
                                                APPLICANT: Wong, Lu Min
APPLICANT: Qian, Xueming
TITLE OF INVENTION: Therapeutic Human Anti-
FILE REFERENCE: 01,1554
CURRENT APPLICATION NUMBER: US/10/656,769
CURRENT FILING DATE: 2003-09-05
NUMBER OF SEQ ID NOS: 79
SOFTWARE: Patentin version 3.0
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Best Local Similarity 96.3%;
Matches 104; Conservative
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APPLICANT: Witte, Alison
APPLICANT: Vezina, Chris
APPLICANT: Wong, Lu Min
APPLICANT: Qian, Xueming
                                Chris
             Alison
                                                                                                                                                                                                                                                                                                                 ; ORGANISM: Homo sapiens
US-10-656-769-12
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                                   Vezina,
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US-10-394-471B-16
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US-10-394-471B-1
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LENGTH: 128
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Best Local S
Matches 104
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TITLE OF IN
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Sequence 48, Application US/10408901

Publication No. US20040023313A1

GENERAL INFORMATION:

APPLICANT: Boyle, William

APPLICANT: Bullot, Robin

APPLICANT: Bliot, Robin

APPLICANT: Bliot, Robin

APPLICANT: Medlock, Eugene

APPLICANT: M
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APPLICANT: Elliot, Robin
APPLICANT: Sullivan, John
APPLICANT: Medlock, Eugene
APPLICANT: Medlock, Eugene
APPLICANT: Martin, Francis
TITLE OF INVENTION: Human Anti-OPGL Neutralizing Antibodies
FILE REFERENCE: MBHB 01-1145-A
CURRENT APPLICATION NUMBER: US/10/408,901
CURRENT FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 76
SOFTWARE: PatentIn version 3.0
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Pred. No. 6.7e-40
3; Mismatches
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Pred. No. 1.3e-39;
3; Mismatches
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; Sequence 12, Application US/10656769
; Publication No. US20040097712A1
; GENERAL INFORMATION:
; APPLICANT: Varnum, Brian
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ilarity 95.4%;
Conservative
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95.4%;
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nes 103; Conser
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LENGTH: 109
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; Mismatches
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; Sequence 9, Application US/10173551
; Publication No. US20030232387A1
; GENERAL INFORMATION:
; APPLICANT: Lu, Chafen
; TITLE OF INVENTION: Antibodies that bind alp
; FILE REFERENCE: 1855.2025-000
; CURRENT APPLICATION NUMBER: US/10/173,551
; CURRENT FILING DATE: 2002-06-14
; NUMBER OF SEQ ID NOS: 70
; SOFTWARE: FastSEQ for Windows Version 4.0
TITLE OF INVENTION: Inhibitors
FILE REFERENCE: MBHB 01-1145-A
CURRENT APPLICATION NUMBER: US/10/408,901
CURRENT FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 76
SOFTWARE: PatentIn version 3.0
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US-10-886-838-4
; Sequence 4, Application US/10886838
; Publication No. US20050008642A1
; GENERAL INFORMATION:
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1 Similarity 94.4%;
102; Conservative
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OTHER INFORMATION: CDR2
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TYPE: PRT
ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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LOCATION: (89)
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APPLICANT: Huang, Haichun
APPLICANT: Huang, Haichun
APPLICANT: Sullivan, John
APPLICANT: Sullivan, John
APPLICANT: Martin, Francis
TITLE OF INVENTION: Human Anti-OPGL Neutralizing Antibodies As Sel
TITLE OF INVENTION: Inhibitors
FILE REFERENCE: MBHB 01-1145-A
CURRENT APPLICATION NUMBER: US/10/408,901
CURRENT FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 76
SOFTWARE: Patentin version 3.0
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APPLICANT: Elliot, Robin
APPLICANT: Sullivan, John
APPLICANT: Medlock, Eugene
APPLICANT: Martin, Francis
TITLE OF INVENTION: Human Anti-OPGL Neutralizing Antibodies As
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               CURRENT APPLICATION NUMBER: US/10/656,769
CURRENT FILING DATE: 2003-09-05
NUMBER OF SEQ ID NOS: 79
SOFTWARE: Patentin version 3.0
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US-10-408-901-32
; Sequence 32, Application US/10408901
; Publication No. US20040023313A1
; GENERAL INFORMATION:
; APPLICANT: Boyle, William
; APPLICANT: Haichun
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Local Similarity 94.4%;
hes 102; Conservative
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US-10-408-901-8
                                                                                                                                                              Query Match
Best Local Similarity
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     REFERENCE:
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LENGTH: 235
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US-10-408-901-8
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Publication No. US20050208596A1

GENERAL INFORMATION:

APPLICANT: Siegel, Donald L.

TITLE OF INVENTION: AUTOANTIBODIES AND INHIBITORS THEREOF

FILE REFERENCE: 053893-5050

CURRENT APPLICATION NUMBER: US/11/021,715

CURRENT PILING DATE: 2004-12-23

PRIOR PILING DATE: 2003-07-03

PRIOR PILING DATE: 2002-07-03

PRIOR FILING DATE: 2002-07-03

PRIOR FILING DATE: 2002-07-03

PRIOR FILING DATE: 2002-09-18

NUMBER OF SEQ ID NOS: 153

SOFTWARE: Patentin version 3.2
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APPLICANT: Hoffmann-La Roche Inc.

TITLE OF INVENTION: Antibodies against insulin-like
TITLE OF INVENTION: thereof
FILE REFERENCE: 21695
CURRENT APPLICATION NUMBER: US/10/886,838
CURRENT FILING DATE: 2004-07-08
PRIOR APPLICATION NUMBER: EP 03015526
PRIOR FILING DATE: 2003-07-10
NUMBER OF SEQ ID NOS: 8
SOFTWARE: PatentIn version 3.2
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Pred. No. 2.2e-39;
); Mismatches 3;
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Pred. No. 1.8e-39;
2; Mismatches 3;
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Matches 103; Conservative
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102; Conservative
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US-10-886-838-4
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US-11-021-715-101
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US-11-021-715-101
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LENGTH: 108
TYPE: PRT
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2: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

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4: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep:*

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19, Appl 24, Appl 1049, Ap 22, Appl 31, Appl 17, Appl 138, Appl 131, Appl 33, Appl 332, Appl 27, Appl 27, Appl 27, Appl 804, App 804, App 806, App 806, App 806, App 806, Appl 807, Appl 1238, Appl 1238, Appl 805, Appl 806, Appl 807, Appl 807, Appl 807, Appl escription Sequence Seq US-11-093-274-23 US-11-093-274-24 US-11-093-274-24 US-11-093-274-24 US-11-093-274-22 US-11-093-274-22 US-11-093-274-22 US-11-093-274-33 US-11-084-554-138 US-11-084-554-131 US-11-084-554-131 US-11-084-554-131 US-11-000-463-332 US-11-000-463-805 US-11-000-463-806 US-11-000-463-806 US-11-000-463-806 US-11-000-463-806 US-11-000-463-806 US-11-056-825-8 US-11-056-825-8 US-11-056-825-4 B L0LLL00LLLLLLCLLLLLLLLL Length Query Match 531.5 530.5 530.5 530.5 520.5 Score

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RESULT 2
US-10-721-763-19
; Sequence 19, Application US/10721763
; Publication No. US20050249729A1
; GENERAL INFORMATION:
; APPLICANT: KIRIN BEER KABUSHIKI KAISHA
; TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY
; FILE REFERENCE: PH-1573-PCT
; CURRENT APPLICATION NUMBER: US/10/721,763
; CURRENT FILING DATE: 2003-11-26
; PRIOR APPLICATION NUMBER: JP2001-150213

RESULT US-10-7

Sequence 1264, Ap Sequence 1219, Ap Sequence 178, App Sequence 178, Appli Sequence 1534, App Sequence 127, Appli Sequence 127, App Sequence 127, App Sequence 130, App Sequence 1035, App Sequence 4, Appli Sequence 4, Appli Sequence 135, App Sequence 135, App Sequence 135, App Sequence 135, App Sequence 115, Appli Sequence 115, Appli Sequence 111, Appli Sequence 161, App Sequence 161, App	USES	9; DB 7; Length 108; tches 2; Indels 0; Gaps 0; VYSYLAWYQQKPGQAPRLLIYDASNRATGIPA 60 
US-11-054-515-1264 US-11-054-515-1219 US-11-054-515-1499 US-11-064-174-178 US-11-054-515-1534 US-11-054-515-905 US-11-054-669-84 US-11-054-515-905 US-11-054-515-1037 US-11-084-554-127 US-11-084-554-127 US-11-064-174-177 US-11-064-174-177 US-11-064-174-177 US-11-064-174-177 US-11-064-174-135 US-11-064-174-115	ALIGNMENTS 093274 1. phine M. ribodies AND THEIR -US1 5/11/093,274 -28 557,741	Score 53 Pred. No 3; Misma TLSCRASOS           TLSCRASOS           PEDFAVYYC
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26 29 29 29 30 30 31 33 34 34 34 35 36 44 44 45 45 46 46 46 47 47 47 47 47 47 47 47 47 47 47 47 47	RESULT 1 US-11-093-274-23 ; Sequence 23, Applicati ; Publication No. US2005 ; GENERAL INFORMATION: ; APPLICANT: Graziano, ; APPLICANT: Cardarell ; APPLICANT: Cardarell ; APPLICANT: Kempe, Th ; APPLICANT: Kempe, Th ; APPLICANT: Srinivasa ; TITLE OF INVENTION: I ; FILE REFERENCE: 04280 ; CURRENT APPLICATION NUM ; FILE REFERENCE: 20 ; CURRENT APPLICATION NUM ; PRIOR FILING DATE: 20 ; NUMBER OF SEQ ID NOS: ; SOFTWARE: PatentIn ve ; SEQ ID NO 23 ; LENGTH: 108 ; TYPE: PRT ; ORGANISM: HOMO Bapie US-11-093-274-23	Query Match         Best Local Sim         Matches 103;         Qy       1 EI         Db       1 EI         Qy       61 RF         Db       61 RF         Db       61 RF

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Publication No. US2005025532A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523P3
CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
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APPLICANT: Cardarelli, Josephine M.
APPLICANT: Kempe, Thomas
APPLICANT: Kempe, Thomas
APPLICANT: Cutter, Beth
APPLICANT: Srinivasan, Mohan
TITLE OF INVENTION: IRTA-5 ANTIBODIES AND THEIR USES
FILE REFERENCE: 04280/1201101-US1
CURRENT APPLICATION NUMBER: US/11/093,274
CURRENT FILING DATE: 2005-03-28
PRIOR APPLICATION NUMBER: 60/557,741
                                                                                                                                                                                                           Score 531.5; DB 6;
Pred. No. 1.9e-34;
; Mismatches 2;
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Pred. No. 2e-34;
; Mismatches 2;
                 JP2001-243040
                              -08-09
R: JP2001-314489
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           PRIOR APPLICATION NUMBER: JP200
PRIOR FILING DATE: 2001-08-09
PRIOR APPLICATION NUMBER: JP200
PRIOR FILING DATE: 2001-10-11
NUMBER OF SEQ ID NOS: 45
SOFTWARE: Patentin Ver. 2.1
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ilarity 96.3%;
Conservative
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llarity 96.3%;
Conservative
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R OF SEQ ID NOS: 41
ARE: Patentin version 3.3
2001-05-18
                                                                                                                                                         ; ORGANISM: Homo sapiens US-10-721-763-19
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US-11-093-274-24
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Best Local Similarity
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Best Local Similarity
FILING DATE:
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                                                                                                            SEQ ID NO 19
LENGTH: 129
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Pred. No. 5e-34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22, Application US/11093274

Publication No. US20050266008A1

GENERAL INFORMATION:
APPLICANT: Graziano, Robert
APPLICANT: Cardarelli, Josephine M.
APPLICANT: Kempe, Thomas
APPLICANT: Cutter, Beth
APPLICANT: Srinivasan, Mohan
TITLE OF INVENTION: IRTA-5 ANTIBODIES AND THI
FILE REFERENCE: 04280/1201101-US1
CURRENT APPLICATION NUMBER: US/11/093,274
CURRENT FILING DATE: 2005-03-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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EQ ID NO 1049
LENGTH: 251
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PRIOR APPLICATION NUMBER: 09/880,748
PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2000-10-17
OR FILING DATE: 2004-02-11
OR APPLICATION NUMBER: 60/580,347
OR FILING DATE: 2004-06-18
OR APPLICATION NUMBER: 10/293,418
OR FILING DATE: 2002-11-14
OR APPLICATION NUMBER: 60/331,469
OR FILING DATE: 2001-11-16
OR APPLICATION NUMBER: 60/340,817
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PRIOR FILING DATE: 2004-03-29
NUMBER OF SEQ ID NOS: 41
SOFTWARE: Patentin version 3.2
EQ ID NO 22
LENGTH: 108
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illarity 95.4%;
Conservative
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Best Local Similarity 92.6%;
Matches 100; Conservative
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Best Local Similarity
Matches 103; Conser
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US-11-040-159-17
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LENGTH: 107
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LENGTH: 95
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neg 3;
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Pred. No. 9.1e-34;
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; Sequence 23, Application US/10721763
; Publication No. US20050249729A1
; Publication No. US20050249729A1
; GENERAL INFORMATION:
; APPLICANT: KIRIN BEER KABUSHIKI KAISHA
; TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY
; FILE REFERENCE: PH-1573-PCT
; CURRENT APPLICATION NUMBER: US/10/721,763
; CURRENT FILING DATE: 2003-11-26
; PRIOR APPLICATION NUMBER: JP2001-150213
; PRIOR FILING DATE: 2001-05-18
; PRIOR FILING DATE: 2001-08-09
; PRIOR FILING DATE: 2001-08-09
; PRIOR APPLICATION NUMBER: JP2001-314489
                                                                                                                                                               APPLICANT: KIRIN BEER KABUSHIKI KAISHA
TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY
FILE REFERENCE: PH-1573-PCT
CURRENT APPLICATION NUMBER: US/10/721,763
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: JP2001-150213
PRIOR FILING DATE: 2001-05-18
PRIOR PILING DATE: 2001-08-09
PRIOR FILING DATE: 2001-08-09
PRIOR FILING DATE: 2001-10-11
NUMBER OF SEQ ID NOS: 45
SOFTWARE: Patentin Ver. 2.1
                                                                                                  US-10-721-763-31; Sequence 31, Application US/10721763; Publication No. US20050249729A1; GENERAL INFORMATION:
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nilarity 95.4%;
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NUMBER OF SEQ ID NOS: 45
SOFTWARE: Patentin Ver. 2.1
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Best Local Similarity
Matches 103; Conser
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ENGTH: 128
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LENGTH: 129
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APPLICANT: Yarranton, Geoffrey T.
APPLICANT: Kalobios, Inc.
TITLE OF INVENTION: Antibody Specificity Transfer Using Minimal Essential
TITLE OF INVENTION: Binding Determinants
FILE REFERENCE: 021167-001730US
CURRENT APPLICATION NUMBER: US/11/040,159
CURRENT FILING DATE: 2005-01-20
PRIOR APPLICATION NUMBER: US 60/537,364
PRIOR FILING DATE: 2004-01-20
PRIOR FILING DATE: 2004-02-23
NUMBER OF SEQ ID NOS: 133
SOFTWARE: Patentin Ver. 2.1
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Pred. No. 3.4e-31;
3; Mismatches 5;
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TITLE OF INVENTION: SUPER HUMANIZED ANTIBODIES
FILE REFERENCE: 30219/US/3
CURRENT APPLICATION NUMBER: US/11/054,669
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 10/194,975
PRIOR FILING DATE: 2002-07-12
PRIOR FILING DATE: 2001-07-12
NUMBER OF SEQ ID NOS: 124
SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 on of Artificial antibody 1F1 VL
                                                                                                                      Sequence 17, Application US/11040159; Publication No. US2005055552A1; GENERAL INFORMATION:
APPLICANT: Flynn, Peter
APPLICANT: Luehrsen, Kenneth
APPLICANT: Balint, Robert F.
APPLICANT: Beblington, Christopher R.
APPLICANT: Bebbington, Christopher R.
APPLICANT: Yarranton, Geoffrey T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-11-054-669-86; Sequence 86, Application US/11054669; Publication No. US20050261480A1; GENERAL INFORMATION:
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; OTHER INFORMATION: anti-PcrV an
US-11-040-159-17
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llarity 89.9%;
Conservative
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Best Local Similarity
Matches 98; Conser
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                                                                                                                                                                                                                                             Sequence 138, Application US/11084554
; Publication No. US20050260679A1
; GENERAL INFORMATION:
; APPLICANT: Kellermann, Sirid-Ai
; APPLICANT: Green, Larry L.
; APPLICANT: Korver, Wouter
; TITLE OF INVENTION: REDUCING THE RISK OF HUMAN ANTI-HUMAN
; TITLE OF INVENTION: ANTIBODIES THROUGH V GENE MANIPULATION
; FILE REFERENCE: ABGENIX.100A
; CURRENT APPLICATION NUMBER: US/11/084,554
; CURRENT FILING DATE: 2005-03-17
; PRIOR APPLICATION NUMBER: 60/554,372
; PRIOR FILING DATE: 2004-03-19
  Length 95;
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 Score 487; DB 7;
Pred. No. 3.4e-31
); Mismatches
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Pred. No. 3.4e-31
); Mismatches
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APPLICANT: Cardarelli, Josephine M.
APPLICANT: Kempe, Thomas
APPLICANT: Cutter, Beth
APPLICANT: Srinivasan, Mohan
TITLE OF INVENTION: IRTA-5 ANTIBODIES AND THI
FILE REFERENCE: 04280/1201101-US1
CURRENT APPLICATION NUMBER: US/11/093,274
CURRENT FILING DATE: 2005-03-28
PRIOR APPLICATION NUMBER: 60/557,741
PRIOR FILING DATE: 2004-03-29
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SOFTWARE: FastSEQ for Windows Version
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 33, Application US/11093274 Publication No. US20050266008A1 GENERAL INFORMATION:
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SOFTWARE: Patentin version
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ORGANISM: Homo sapiens
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APPLICANT: Green, Larry L.

APPLICANT: Green, Larry L.

APPLICANT: Korver, Wouter

TITLE OF INVENTION: ANTIBODIES THROUGH V GENE MANIPULATION

TITLE OF INVENTION: ANTIBODIES THROUGH V GENE MANIPULATION

FILE REFERENCE: ABGENIX.100A

CURRENT APPLICATION NUMBER: 105/11/084,554

CURRENT FILING DATE: 2005-03-17

PRIOR FILING DATE: 2004-03-19

PRIOR FILING DATE: 2004-05-24

NUMBER OF SEQ ID NOS: 266

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 131

LENGTH: 95

TYPE: PRT

ORGANISM: Homo sapiens
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Pred. No. 5.5e-30;
0; Mismatches 3
                              Score 480; DB 7;
Pred. No. 1.1e-30;
); Mismatches 1
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TITLE OF INVENTION: SUPER HUMANIZED ANTIBODIES
FILE REFERENCE: 30219/US/3
CURRENT APPLICATION NUMBER: US/11/054,669
CURRENT FILING DATE: 2005-02-08
PRIOR APPLICATION NUMBER: US 10/194,975
PRIOR FILING DATE: 2002-07-12
PRIOR PILING DATE: 2001-07-12
PRIOR FILING DATE: 2001-07-12
NUMBER OF SEQ ID NOS: 124
SOFTWARE: Patentin version 3.3
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Publication No. US20050260679A1
GENERAL INFORMATION:
APPLICANT: Kellermann, Sirid-Ai
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ilarity 98.9%;
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US-11-093-274-33
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ENGTH: 95
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                            Length 95;
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APPLICANT: Wang, Zhiwei
APPLICANT: Wehrman, Tom
APPLICANT: Zhang, Jie
APPLICANT: Zhou, Ping
APPLICANT: Cao, Yi-Cheng
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides
FILE REFERENCE: 785CIP4CN
CURRENT APPLICATION NUMBER: US/11/000,463
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Pred. No. 2.6e-29;
; Mismatches 10;
                          Score 471; DB 7; Pred. No. 5.5e-30; ); Mismatches 3;
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                                                                                                                                                                                                                                                             Sequence 27, Application US/10721763

Publication No. US20050249729A1

GENERAL INFORMATION:

APPLICANT: KIRIN BEER KABUSHIKI KAISHA

TITLE OF INVENTION: ANTI TRAIL-R ANTIBODY

FILE REFERENCE: PH-1573-PCT

CURRENT APPLICATION NUMBER: US/10/721,763

CURRENT FILING DATE: 2003-11-26

PRIOR APPLICATION NUMBER: JP2001-150213

PRIOR FILING DATE: 2001-05-18

PRIOR FILING DATE: 2001-08-09

PRIOR FILING DATE: 2001-08-09

PRIOR FILING DATE: 2001-10-11

NUMBER OF SEQ ID NOS: 45

SOFTWARE: PATENTIN Ver. 2.1
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; Sequence 332, Application US/11000463
; Publication No. US20050266423A1
; GENERAL INFORMATION:
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                          th 82.9%;
Similarity 96.8%;
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Liu, Chenghua
Asundi, Vinod
Chen, Rui-hong
Qian, Xiaohong E
Wang, Zhiwei
Wehrman, Tom
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US-11-084-554-131
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SEQ ID NO 2
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         PRIOR APPLICATION NUMBER: PCT/USO1/02623
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 09/922,279
PRIOR FILING DATE: 2001-08-03
PRIOR FILING DATE: 2000-01-25
PRIOR APPLICATION NUMBER: 09/491,404
PRIOR FILING DATE: 2000-07-17
PRIOR FILING DATE: 2000-07-17
PRIOR FILING DATE: 2000-09-03
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-09-15
NUMBER OF SEQ ID NOS: 944
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 332
LENGTH: 307
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Published\_Applications Nucleic Acid and Published\_Applications Amino Acid database searches now generate two sets of results each. The Published\_Applications databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Newly published applications will appear in the Published Applications. New databases: older published applications make up the Published Applications make up the Published Applications as a second at a second and the Published Applications are not the Published Applications as a second at a second and the Published Applications are not the Published Applications as a second at a second and the Published Applications are not the Published Applications as a second at a second and the Published Applications are not the Published Applications as a second at a second and the Published Applications are not the Published Applications and the Published Applications are not the Pu

Searches run against Nucleic Acid Published Applications produce two sets of results, with the extensions rupbm (Published Applications NA New).

Searches run against Amino Acid Published Applications produce two sets of results, with the extensions Searches run against Amino Acid Published Applications AA New).

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Published Applications AA New:\*

1: /cgn2\_6/ptodata/1/pubpaa/US08\_NEW\_PUB.pep:\*
2: /cgn2\_6/ptodata/1/pubpaa/US07\_NEW\_PUB.pep:\*
3: /cgn2\_6/ptodata/1/pubpaa/US07\_NEW\_PUB.pep:\*
4: /cgn2\_6/ptodata/1/pubpaa/US09\_NEW\_PUB.pep:\*
5: /cgn2\_6/ptodata/1/pubpaa/US09\_NEW\_PUB.pep:\*
6: /cgn2\_6/ptodata/1/pubpaa/US10\_NEW\_PUB.pep:\*
7: /cgn2\_6/ptodata/1/pubpaa/US11\_NEW\_PUB.pep:\*
8: /cgn2\_6/ptodata/1/pubpaa/US11\_NEW\_PUB.pep:\*

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Post-processing: Minimum Match Maximum Match Listing first Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Sequence 1731, Ap	26,	512,	153,	25, 7	Sequence 1330, Ap	e 839,	Sequence 1394, Ap	981,	140,	Sequence 8, Appli	397,	110	1115	1724	1725	14,	911,	1113	1119	1627	Sequence 1117, Ap	1118	924,	Sequence 892, App
SUMMARIES	ID	US-11-054-515-1731	US-11-064-174-26	US-11-054-515-512	US-11-064-174-153	US-11-064-174-25	US-11-054-515-1330	US-11-054-515-839	US-11-054-515-1394	US-11-054-515-981	US-11-064-174-140	US-11-084-554-8	US-11-054-515-397	US-11-054-515-1102	US-11-054-515-1115	US-11-054-515-1724	US-11-054-515-1725	US-11-040-159-14	US-11-054-515-911	US-11-054-515-1113	US-11-054-515-1119	US-11-054-515-1627	US-11-054-515-1117	US-11-054-515-1118	US-11-054-515-924	US-11-054-515-892
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Sequence 1105, Ap Sequence 1108, Ap	1120, A	13, App	152, Ap	Sequence 896, App	9, Appl	18,	146, A	Sequence 1111, Ap	17, Ap	983, A	<u>,</u>	Sequence 1519, Ap	Sequence 16, Appl	e, A	881, A	Sequence 1109, Ap	Sequence 1421, Ap	Sequence 5, Appli
US-11-054-515-1105 US-11-054-515-1108	15-11	US-11-064-174-13	US-11-064-174-152	US-11-054-515-896	US-11-127-677-9	US-11-155-775-18	US-11-064-174-146	US-11-054-515-1111	US-11-064-174-17	US-11-054-515-983	US-11-054-515-1110	US-11-054-515-1519	US-11-040-159-16	US-11-054-515-956	US-11-054-515-881	US-11-054-515-1109	US-11-054-515-1421	US-11-054-515-5
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                                                                                            APPLICANT: Ruben et al.

APPLICANT: Ruben et al.

TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523P3

CURRENT APPLICATION NUMBER: US/11/054,515

CURRENT FILING DATE: 2005-02-10

PRIOR PILING DATE: 2004-06-18

PRIOR PILING DATE: 2004-06-18

PRIOR FILING DATE: 2004-06-18

PRIOR FILING DATE: 2002-11-14

PRIOR FILING DATE: 2001-11-16

PRIOR FILING DATE: 2001-11-16

PRIOR FILING DATE: 2001-12-19

PRIOR FILING DATE: 2001-06-15

PRIOR FILING DATE: 2001-06-15

PRIOR FILING DATE: 2001-06-15

PRIOR FILING DATE: 2001-06-15

PRIOR PILING DATE: 2001-06-25

PRIOR PILING DATE: 2001-03-21

PRIOR PILING DATE: 2001-03-21

PRIOR PILING DATE: 2001-03-16

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PRIOR FILING DATE: 2000-10-17
Remaining Prior Application data removed
NUMBER OF SEQ ID NOS: 3247
LENGTH: 252
                                  Sequence 1731, Application US/11054515
Publication No. US20050255532A1
GENERAL INFORMATION:
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Best Local Similarity 86.9%;
Matches 113; Conservative
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 105; Conser
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                                                                                                                                                    Sequence 26, Application US/11064174;
Sequence 26, Application US/11064174;
Publication No. US20050282252A1

GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/11/064,174

CURRENT APPLICATION NUMBER: US/09/240,274

PRIOR PILING DATE: 1999-01-29
PRIOR FILING DATE: 1999-01-29
PRIOR FILING DATE: 1998-04-10
PRIOR PPLICATION NUMBER: 60/028,550
PRIOR APPLICATION NUMBER: 60/028,550
PRIOR APPLICATION NUMBER: 60/028,550
PRIOR APPLICATION NUMBER: 2005-02-10-11
SOFTWARE: Patentin Ver. 2.0
ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRLEYYDILT-GYYYYYGMDVW
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APPLICANT: Ruben et al.

TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS

FILE REFERENCE: PF523P3

CURRENT APPLICATION NUMBER: US/11/054,515

CURRENT APPLICATION NUMBER: 60/543,296

PRIOR FILING DATE: 2004-02-11

PRIOR FILING DATE: 2004-06-18

PRIOR FILING DATE: 2002-11-14

PRIOR APPLICATION NUMBER: 60/331,469

PRIOR FILING DATE: 2001-11-16

PRIOR FILING DATE: 2001-12-19

PRIOR FILING DATE: 2001-06-15
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Pred. No. 1.2e-41
3; Mismatches 13
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1-064-174-26
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Best Local Similarity 84.1%;
Matches 106; Conservative
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US-11-054-515-512
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ENGTH: 126
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ATTLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/11/064,174
CURRENT FILING DATE: 2005-02-22
PRIOR APPLICATION NUMBER: G0/081,380
PRIOR FILING DATE: 1999-04-10
PRIOR FILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 153
LENGTH: 126
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US-11-064-174-153
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Pred. No. 6.1e-41
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Pred. No. 4e-41;
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            PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2000-10-17
Remaining Prior Application data removed - NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 512
LENGTH: 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 153, Application US/11064174 Publication No. US20050282252A1 GENERAL INFORMATION:
60/293,499
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523P3
CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT PILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR PILING DATE: 2004-02-11
PRIOR PILING DATE: 2004-06-18
PRIOR PILING DATE: 2004-06-18
PRIOR PILING DATE: 2001-11-14
PRIOR PILING DATE: 2001-11-16
PRIOR PILING DATE: 2001-12-19
PRIOR PILING DATE: 2001-12-19
PRIOR PILING DATE: 2001-06-15
                                                                                                                             APPLICANT: Siegel, Donald L.

TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATE
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT APPLICATION NUMBER: US/11/064,174
CURRENT PILING DATE: 2005-02-22
PRIOR APPLICATION NUMBER: US/09/240,274
PRIOR PILING DATE: 1999-01-29
PRIOR PILING DATE: 1996-04-10
PRIOR PILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
SOFTWARE: PATENTI VET: 2.0
SEQ ID NO 25
LENGTH: 126
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13;
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Pred. No. 5.8e-41;
; Mismatches 13
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; Sequence 1330, Application US/11054515
; Publication No. US20050255532A1
; GENERAL INFORMATION:
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                                                                                      Sequence 25, Application US/11064174 Publication No. US20050282252A1 GENERAL INFORMATION:
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Best Local Similarity 83.3%;
Matches 105; Conservative
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FILING DATE: 2001-05-
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ORGANISM: Homo sapiens
121 TVTVSS 126
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                                                                                  See File Wrapper
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CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
PRIOR FILING DATE: 2004-06-18
PRIOR APPLICATION NUMBER: 10/293,418
PRIOR FILING DATE: 2002-11-14
PRIOR FILING DATE: 2002-11-14
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
               PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
Remaining Prior Application data removed -
NUMBER OF SEQ ID NOS: 3247
SEQ ID NO 1330
LENGTH: 247
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Pred. No.
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1-03-21
ER: 60/276,248
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Pred.
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o. US2005025532A1
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TITLE OF INVENTION: Antibodies that
FILE REFERENCE: PF523P3
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85.8%;
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APPLICATION NUMBER: 09/8
FILING DATE: 2001-06-15
APPLICATION NUMBER: 60/3
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84.5%;
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0 839
: 256
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R APPLICATION NUMBER: R
FILING DATE: 2001-03-
R APPLICATION NUMBER:
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US-11-054-515-1330
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109; Conserv
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Sequence 839, Applica
Publication No. US200
GENERAL INFORMATION:
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109; Conser
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ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRG---IAAGGNYYYYGMDVWG
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Pred. No. 3.9e-40;
; Mismatches 13;
                                                                                                                                                                                                                                                                                                                FILE REFERENCE: PF523P3
CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT PILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
PRIOR FILING DATE: 2004-06-18
PRIOR FILING DATE: 2002-11-14
PRIOR FILING DATE: 2001-11-16
PRIOR FILING DATE: 2001-11-16
PRIOR PPLICATION NUMBER: 60/340,817
PRIOR FILING DATE: 2001-12-19
PRIOR FILING DATE: 2001-06-15
PRIOR FILING DATE: 2001-05-25
PRIOR FILING DATE: 2001-05-25
PRIOR FILING DATE: 2001-05-25
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2000-10-17
PRIOR FILING DATE: 2000-10-17
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3-11-054-515-981
Sequence 981, Application US/11054515
Publication No. US20050255532A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
                                                                                                                                                                                                        US-11-054-515-1394
Sequence 1394, Application US/1105451
Publication No. US20050255532A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that
FILE REFERENCE: PF523P3
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illarity 82.2%;
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Best Local Similarity
Matches 106; Conser
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US-11-054-515-1394
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APPLICANT: Siegel, Donald L.

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

FILE REFERENCE: 09596-42U2

CURRENT APPLICATION NUMBER: US/11/064,174

CURRENT FILING DATE: 2005-02-22

PRIOR APPLICATION NUMBER: US/09/240,274

PRIOR FILING DATE: 1999-01-29

PRIOR FILING DATE: 1998-04-10

PRIOR FILING DATE: 1996-10-11

NUMBER OF SEQ ID NOS: 224

SOFTWARE: Patentin Ver. 2.0

LENGTH: 125
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Antibodies that Immunospecifically Bind BLyS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 254;
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Pred. No. 5.3e-40
; Mismatches 1
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                                     MBER: US/11/054,515
2005-02-10
                                                                                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
Remaining Prior Application data removed
NUMBER OF SEQ ID NOS: 3247
                               CURRENT APPLICATION NUMBER: US/11/054, ECURRENT APPLICATION NUMBER: US/11/054, ECURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543, 296
PRIOR FILING DATE: 2004-02-11
PRIOR APPLICATION NUMBER: 60/580, 347
PRIOR FILING DATE: 2002-11-14
PRIOR FILING DATE: 2001-11-16
PRIOR PILING DATE: 2001-11-16
PRIOR PILING DATE: 2001-11-16
PRIOR PILING DATE: 2001-12-19
PRIOR PILING DATE: 2001-12-19
PRIOR PILING DATE: 2001-06-15
PRIOR FILING DATE: 2001-06-15
PRIOR PILING DATE: 2001-06-15
PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/277,379
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84.4%;
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Best Local Similarity
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US-11-064-174-140
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REPERENCE: PF523P3
CURRENT APPLICATION NUMBER: US/11/054,515
CURRENT FILING DATE: 2005-02-10
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                                                                                                     Length 125;
                                       Length 125;
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APPLICANT: Green, Larry L.
TITLE OF INVENTION: REDUCING THE RISK OF HUMAN ANTI-HUMAN
TITLE OF INVENTION: ANTIBODIES THROUGH V GENE MANIPULATION
FILE REFERENCE: ABGENIX.100A
                                                               Indels
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Pred. No. 3.6e-40;
1; Mismatches 13;
                                                            11;
                                    Score 548.5; DB 7;
Pred. No. 3e-40;
); Mismatches 11;
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CURRENT FILING DATE: 2005-03-17
PRIOR APPLICATION NUMBER: 60/554,372
PRIOR FILING DATE: 2004-03-19
PRIOR PILING DATE: 2004-05-24
NUMBER OF SEQ ID NOS: 266
SOFTWARE: FastSEQ for Windows Version 4.0
antibody
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Publication No. US20050260679A1
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109; Conservative
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Best Local Similarity 83.3%;
Matches 105; Conservative
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APPLICANT: Green, Lari
APPLICANT: Korver, Wol
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TVTVSS 125
; OTHER INFORMATION:
US-11-064-174-140
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523P3
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ed. No. 9.2e-40;
Mismatches 17
PRIOR APPLICATION NUMBER: 60/580,347
PRIOR FILING DATE: 2004-06-18
PRIOR FILING DATE: 2004-06-18
PRIOR PILING DATE: 2002-11-14
PRIOR PILING DATE: 2002-11-14
PRIOR PILING DATE: 2001-11-16
PRIOR FILING DATE: 2001-11-16
PRIOR FILING DATE: 2001-12-19
PRIOR FILING DATE: 2001-06-15
PRIOR PPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-06-15
PRIOR PPLICATION NUMBER: 60/293,499
PRIOR PPLICATION NUMBER: 60/293,499
PRIOR PPLING DATE: 2001-05-25
PRIOR PPLING DATE: 2001-03-21
PRIOR PPLING DATE: 2001-03-21
PRIOR PPLING DATE: 2001-03-16
PRIOR PILING DATE: 2001-03-16
PRIOR PILING DATE: 2000-10-17
Remaining Prior Application data removed - 8
NUMBER OF SEQ ID NOS: 3247
TYPE: PPT
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CURRENT FILING DATE: 2005-02-10
PRIOR APPLICATION NUMBER: 60/543,296
PRIOR FILING DATE: 2004-02-11
PRIOR FILING DATE: 2004-06-18
PRIOR FILING DATE: 2002-11-14
PRIOR FILING DATE: 2002-11-14
PRIOR FILING DATE: 2001-11-16
PRIOR FILING DATE: 2001-11-16
PRIOR FILING DATE: 2001-11-16
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PRIOR FILING DATE: 2001-12-19
PRIOR FILING DATE: 2001-12-19
PRIOR FILING DATE: 2001-06-15
PRIOR FILING DATE: 2001-06-15
PRIOR FILING DATE: 2001-06-15
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Sequence 1102, Application US/11054515
Publication No. US20050255532A1
GENERAL INFORMATION:
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Pred.
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2001-03-21
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APPLICATION NUMBER:
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Pred. No. 9.2e-40;
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Pred. No. 9.2e-40
1; Mismatches 1
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TITLE OF INVENTION: Antibodies that Immunosp.
FILE REFERENCE: PF523P3

CURRENT APPLICATION NUMBER: US/11/054,515

CURRENT FILING DATE: 2005-02-10

PRIOR APPLICATION NUMBER: 60/543,296

PRIOR FILING DATE: 2004-02-11

PRIOR FILING DATE: 2004-06-18

PRIOR FILING DATE: 2002-11-14

PRIOR FILING DATE: 2002-11-14

PRIOR FILING DATE: 2001-11-16

PRIOR FILING DATE: 2001-11-16

PRIOR FILING DATE: 2001-11-16

PRIOR FILING DATE: 2001-11-16

PRIOR FILING DATE: 2001-11-16
PRIOR FILING DATE: 2000-10-17
Remaining Prior Application data removed
NUMBER OF SEQ ID NOS: 3247
EQ ID NO 1102
LENGTH: 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
Remaining Prior Application data removed
NUMBER OF SEQ ID NOS: 3247
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APPLICATION NUMBER: 09/E
FILING DATE: 2001-06-15
APPLICATION NUMBER: 60/2
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FILING DATE: 2001-03-21
APPLICATION NUMBER: 60/
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Matches 106; Conser
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Best Local Similarity
Matches 106; Conser
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US-11-054-515-1115
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GENERAL INFORMATION:

APPLICANT: Ruben et al.

TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523P3

CURRENT APPLICATION NUMBER: US/11/054,515

CURRENT APPLICATION NUMBER: 00/543,296

PRIOR APPLICATION NUMBER: 60/580,347

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Remaining Prior Application data removed
NUMBER OF SEQ ID NOS: 3247
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; Sequence 1724, Application US/11054515
; Publication No. US20050255532A1
; GENERAL INFORMATION:
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Seharaseyon; Jegatheesan----

Sent:

Tuesday, December 27, 2005 3:25 PM

To:

STIC-Biotech/ChemLib

Subject:

Re:09/920137

Importance:

High

Hi,

Please search SEQ ID NO: 7 and 8 of 09/920137 in both the commercial and interference databases.

Thanks in advance, Seyon.

J.Seharaseyon Art Unit 1647 Remsen 4C61 Mailbox 4C70

Phone: (571)-272-0892 Fax: (571)-273-0892 M

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searcher Phone:	NA# AA#:
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